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1. Document ID: US 20020013257 A1

L1: Entry 1 of 2

File: PGPB

Jan 31, 2002

PGPUB-DOCUMENT-NUMBER: 20020013257

PGPUB-FILING-TYPE: new

DOCUMENT-IDENTIFIER: US 20020013257 A1

TITLE: Use of ligands to GABAB receptors

PUBLICATION-DATE: January 31, 2002

INVENTOR-INFORMATION:

NAME

CITY

STATE

COUNTRY

RULE-47

Bernasconi, Raymond

Oberwil

CH

Otten, Uwe

Basel

CH

US-CL-CURRENT: 514/1

Full Title Citation Front Review Classification Date Reference Sequences Affachments

KMMC Draw Desc Image

2. Document ID: US 20010023289 A1

L1: Entry 2 of 2

File: PGPB

Sep 20, 2001

PGPUB-DOCUMENT-NUMBER: 20010023289

PGPUB-FILING-TYPE: new

DOCUMENT-IDENTIFIER: US 20010023289 A1

TITLE: DNA encoding a GABABR2 polypeptide and uses thereof

PUBLICATION-DATE: September 20, 2001

INVENTOR-INFORMATION:

NAME

CITY

STATE

COUNTRY

RULE-47

Jones, Kenneth A.

Bergenfield

NJ

HC

Jones, Reimech A

50150....

MO

US

Laz, Thomas M. Borowsky, Beth

Parlin Montclair NJ NJ US US

US-CL-CURRENT: 536/23.5

Full Title Citation Front Review Classification Date Reference Sequences Attachments

EWMC Draw Desc Image

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Term	Documents
GABAB\$3	0
GABAB.USPT,PGPB.	16
GABABA.USPT,PGPB.	1
GABABLA.USPT,PGPB.	1
GABABRL.USPT,PGPB.	1
GABABRLA.USPT,PGPB.	1
GABABRLB.USPT,PGPB.	1
GABABR2.USPT,PGPB.	1
GABABR2:.USPT,PGPB.	1
GABAB1A.USPT,PGPB.	1
GABAB1AA.USPT,PGPB.	1
(GABAB\$3.AB. OR GABAB\$.TI.).USPT,PGPB.	2

There are more results than shown above. Click here to view the entire set.

Display Format: REV Change Format

Previous Page Next Page

```
GBR2 RAT
                         STANDARD;
                                         PRT;
                                                940 AA.
         GBR2 RAT
     ID
         088871; Q9QWU2; Q9JK36;
    AC
          20-AUG-2001 (Rel. 40, Created)
     DT
5
          20-AUG-2001 (Rel. 40, Last sequence update)
     DT
          20-AUG-2001 (Rel. 40, Last annotation update)
     DT
          GAMMA-AMINOBUTYRIC ACID TYPE B RECEPTOR, SUBUNIT 2 PRECURSOR (GABA-B
     DE
          RECEPTOR 2) (GABA-B-R2) (GB2) (GABABR2).
     DΕ
     GN
          GABBR2.
          Rattus norvegicus (Rat).
10
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          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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          NCBI TaxID=10116;
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     RN
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          Yao W.-J., Johnson M., Gunwaldsen C.A., Huang L.-Y., Tang C., Shen Q.,
     RA
          Salon J.A., Morse K., Laz T., Smith K.E., Nagarathnam D., Noble S.A.,
20
     RA
          Branchek T.A., Gerald C.;
     RA
          "GABA(B) receptors function as a heteromeric assembly of the subunits
     RT
          GABA(B)R1 and GABA(B)R2.";
     RT
          Nature 396:674-679(1998).
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     RA
          Mosbacher J., Bischoff S., Kulik A., Shigemoto R., Karschin A.,
30
     RA
          Bettler B.;
     RA
          "GABA-B receptor subtypes assemble into functional heteromeric
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          complexes.";
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          Nature 396:683-687(1998).
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          "Distribution of the GABA(B) receptor subunit gb2 in rat CNS.";
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          Brain Res. 860:41-52(2000).
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     RN
          [4]
          SEQUENCE FROM N.A.
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          TISSUE=Hypothalamus;
     RC
          Borowsky B., Laz T., Gerald C.;
45
     RA
          Submitted (JAN-1999) to the EMBL/GenBank/DDBJ databases.
     RL
     RN
          [5]
          R1A-R2 INTERACTION, AND VARIANTS TYR-337 AND PRO-19 INS.
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          TISSUE=Hippocampus;
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     RX
          Kuner R., Koehr G., Gruenewald S., Eisenhardt G., Bach A.,
     RA
     RA
          Kornau H.-C.;
          "Role of Heteromer Formation in GABA-B Receptor Function.";
     RT
          Science 283:74-77(1999).
     RL
          -!- FUNCTION: RECEPTOR FOR GABA. THE ACTIVITY OF THIS RECEPTOR IS
55
     CC
              MEDIATED BY G-PROTEINS THAT INHIBITS ADENYLYL CYCLASE ACTIVITY,
     CC
              STIMULATES PHOSPHOLIPASE A2, ACTIVATES POTASSIUM CHANNELS,
     CC
```

INACTIVATES VOLTAGE-DEPENDENT CALCIUM-CHANNELS AND MODULATES INOSITOL PHOSPHOLIPIDS HYDROLYSIS. PLAYS A CRITICAL ROLE IN THE CC FINE-TUNING OF INHIBITORY SYNAPTIC TRANSMISSION. PRE-SYNAPTIC CC GABA-B-R INHIBIT NEUROTRANSMITTER RELEASE BY DOWN-REGULATING CC HIGH-VOLTAGE ACTIVATED CALCIUM CHANNELS, WHEREAS POSTSYNAPTIC 5 CCGABA-B-R DECREASE NEURONAL EXCITABILITY BY ACTIVATING A PROMINENT CC INWARDLY RECTIFYING POTASSIUM (KIR) CONDUCTANCE THAT UNDERLIES THE CC LATE INHIBITORY POSTSYNAPTIC POTENTIALS. NOT ONLY IMPLICATED IN CCSYNAPTIC INHIBITION BUT ALSO IN HIPPOCAMPAL LONG-TERM CC POTENTIATION, SLOW WAVE SLEEP, MUSCLE RELAXATION AND 10 CC ANTINOCICEPTION. CC -!- SUBUNIT: HETERODIMER OF GABA-B-R1 AND GABA-B-R2. NEITHER OF WHICH CC IS EFFECTIVE ON ITS OWN AND HOMODIMERIC ASSEMBLY DOES NOT SEEM TO CC HAPPEN. CC-!- SUBCELLULAR LOCATION: INTEGRAL MEMBRANE PROTEIN. MOREOVER 15 CCCOEXPRESSION OF GABA-B-R1 AND GABA-B-R2 APPEARS TO BE A CCPREREQUISITE FOR MATURATION AND TRANSPORT OF GABA-B-R1 TO THE CC PLASMA MEMBRANE. CC-!- TISSUE SPECIFICITY: EXPRESSED IN BRAIN CORTEX, HIPPOCAMPUS, MEDIAL CC HABENULA, THALAMUS AND CEREBELLUM. COEXPRESSION IS SEEN IN 20 CC CC CEREBELLUM. -!- DEVELOPMENTAL STAGE: ABUNDANT IN BRAIN CORTEX AND CEREBELLUM CCTHROUGHOUT POSTNATAL DEVELOPMENT WHEREAS ITS EXPRESSION IN SPINAL CCCC CORD GRADUALLY DECREASES. -!- DOMAIN: ALPHA-HELICAL PARTS OF THE C-TERMINAL INTRACELLULAR REGION 25 CC MEDIATE HETERODIMERIC INTERACTION WITH GABA-B RECEPTOR 1. CC -!- SIMILARITY: BELONGS TO FAMILY 3 OF G-PROTEIN COUPLED RECEPTORS. CC CC GABA-B RECEPTOR SUBFAMILY. CC30 This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation CC the European Bioinformatics Institute. There are no restrictions on 35 CC its use by non-profit institutions as long as its content is in no CC way modified and this statement is not removed. Usage by and for CC 40 commercial entities requires a license agreement (See http://www.isb-CC sib.ch/announce/ CC or send an email to license@isb-sib.ch). CC 45 DR EMBL; AF074482; AAD03335.1; -. DR EMBL; AJ011318; CAA09592.1; -. DR EMBL; AF058795; AAC63994.1; -. DR EMBL; AF109405; AAD03338.1; -. InterPro; IPR001828; ANF receptor. 50 DR DR InterPro; IPR000337; GPCR Mgr. DR Pfam; PF00003; 7tm 3; 1. Pfam; PF01094; ANF receptor; 1. DR PRINTS; PR00248; GPCRMGR. DR DR PRINTS; PR01176; GABABRECEPTR. 55 DR PRINTS; PR01177; GABAB1RECPTR. PRINTS; PR01178; GABAB2RECPTR. DR

```
PROSITE; PS50099; PRO RICH; 1.
    DR
         PROSITE; PS00979; G PROTEIN RECEP F3_1; FALSE_NEG.
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         PROSITE; PS00980; G PROTEIN RECEP F3 2; FALSE NEG.
    DR
         PROSITE; PS00981; G PROTEIN RECEP F3 3; FALSE NEG.
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    KW
         Postsynaptic membrane; Coiled coil; Polymorphism.
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                      1
                            40
                                    POTENTIAL.
    FT
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    FT
                                    SUBUNIT 2.
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    FT
                    483
                           503
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                           521
                                    CYTOPLASMIC (POTENTIAL).
                    504
    FT
         DOMAIN
         TRANSMEM
                    522
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                                    II (POTENTIAL).
    FT
15
                    543
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                    618
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                                    VI (POTENTIAL).
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    FT
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                                    VII (POTENTIAL).
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25
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         DOMAIN
                    741
                           940
    FT
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    FT
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    FT
                           297
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                            19
    FT
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                     19
                                    P -> PP.
    FT
         VARIANT
                     19
                            19
                    337
                           337
                                    F \rightarrow Y.
    FT
         VARIANT
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35
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    Qу
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    Qу
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    Qу
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	Qу		RCLRKNLLAAMEGYIGVDFEPLSSKQIKTISGKTPQQYEREYNNKRSGVGPSKFHGYAYD	360
10	Db	300	RCLRRSLLAAMEGYIGVDFEPLSSKQIKTISGKTPQQFEREYNSKRSGVGPSKFHGYAYD	359
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	Db	360	GIWVIAKTLQRAMETLHASSRHQRIQDFNYTDHTLGKIILNAMNETNFFGVTGQVVFRNG	419
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20	Qy		LYSILSALTILGMIMASAFLFFNIKNRNQKLIKMSSPYMNNLIILGGMLSYASIFLFGLD	540
	Db	480	LYSILSALTILGMIMASAFLFFNIKNRNQKLIKMSSPYMNNLIILGGMLSYASIFLFGLD	539
	Qу		GSFVSEKTFETLCTVRTWILTVGYTTAFGAMFAKTWRVHAIFKNVKMKKKIIKDQKLLVI	600
25	Db	540	GSFVSEKTFETLCTVRTWILTVGYTTAFGAMFAKTWRVHAIFKNVKMKKKIIKDQKLLVI	599
	QУ		VGGMLLIDLCILICWQAVDPLRRTVEKYSMEPDPAGRDISIRPLLEHCENTHMTIWLGIV	660
20	Db	600	VGGMLLIDLCILICWQAVDPLRRTVERYSMEPDPAGRDISIRPLLEHCENTHMTIWLGIV	659
30	QУ		YAYKGLLMLFGCFLAWETRNVSIPALNDSKYIGMSVYNVGIMCIIGAAVSFLTRDQPNVQ	720
	Db	660	YAYKGLLMLFGCFLAWETRNVSIPALNDSKYIGMSVYNVGIMCIIGAAVSFLTRDQPNVQ	719
35	QУ		FCIVALVIIFCSTITLCLVFVPKLITLRTNPDAATQNRRFQFTQNQKKEDSKTSTSVTSV	
	Db	720	FCIVALVIIFCSTITLCLVFVPKLITLRTNPDAATQNRRFQFTQNQKKEDSKTSTSVTSV	779
40	QУ		NQASTSRLEGLQSENHRLRMKITELDKDLEEVTMQLQDTPEKTTYIKQNHYQELNDILNL	
	Db	780	NQASTSRLEGLQSENHRLRMKITELDKDLEEVTMQLQDTPEKTTYIKQNHYQELNDILSL	839
	Qу		GNFTESTDGGKAILKNHLDQNPQLQWNTTEPSRTCKDPIEDINSPEHIQRRLSLQLPILH	
45	Db	840	GNFTESTDGGKAILKNHLDQNPQLQWNTTEPSRTCKDPIEDINSPEHIQRRLSLQLPILH	899
	Qу	901	HAYLPSIGGVDASCVSPCVSPTASPRHRHVPPSFRVMVSGL 941	
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Application/Control Number: 09/601,582

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GBR2 HUMAN
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                         STANDARD;
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          075899; 075974; 075975; Q9UNS9; Q9UNR1; Q9P1R2;
     AC
          20-AUG-2001 (Rel. 40, Created)
     DT
          20-AUG-2001 (Rel. 40, Last sequence update)
 5
     \mathsf{D}\mathbf{T}
          20-AUG-2001 (Rel. 40, Last annotation update)
     DT
          GAMMA-AMINOBUTYRIC ACID TYPE B RECEPTOR, SUBUNIT 2 PRECURSOR (GABA-B
     DΕ
          RECEPTOR 2) (GABA-B-R2) (GB2) (GABABR2) (G PROTEIN-COUPLED RECEPTOR
     DE
          51) (GPR 51) (HG20).
     DE
10
     GN
          GABBR2 OR GPR51.
          Homo sapiens (Human).
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          Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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          Barnes A.A., Emson P., Foord S.M., Marshall F.H.;
20
     RA
          "Heterodimerization is required for the formation of a functional
     RT
          GABA(B) receptor.";
     RT
          Nature 396:679-682(1998).
     RL
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          PARTIAL SEQUENCE FROM N.A. (ISOFORMS 2A; 2B AND 2C).
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          Clark J.A., Mezey E., Lam A.S., Bonner T.I.;
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          "Distribution of the GABA(B) receptor subunit gb2 in rat CNS.";
     RT
          Brain Res. 860:41-52(2000).
30
     RL
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          SEQUENCE FROM N.A. (ISOFORM 2A).
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          Liu M., Parker R., McCrea K., Watson J., Baker E., Sutherland G.,
     RA
     RA
          Herzog H.;
          "Cloning and characterization of a novel human GABA-B receptor subtype
35
     RT
          with high affinity for GABA and low affinity for baclofen.";
     RT
          Submitted (NOV-1998) to the EMBL/GenBank/DDBJ databases.
     RL
     RN
          SEQUENCE FROM N.A. (ISOFORM 2A).
     RP
40
          TISSUE=Hippocampus;
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          Borowsky B., Laz T., Gerald C.;
     RA
          Submitted (JAN-1999) to the EMBL/GenBank/DDBJ databases.
     RL
     RN
          SEQUENCE FROM N.A. (ISOFORM 2A).
     RP
45
          TISSUE=Fetal brain;
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          MEDLINE=99189236; PubMed=10087195;
     RX
          Ng G.Y.K., McDonald T., Bonnert T., Rigby M., Heavens R., Whiting P.,
     RA
          Chateauneuf A., Coulombe N., Kargman S., Caskey T., Evans J.F.,
     RA
          O'Neill G.P., Liu Q.;
     RA
          "Cloning of a novel G-protein-coupled receptor GPR 51 resembling GABAB
50
     RT
          receptors expressed predominantly in nervous tissues and mapped
     RT
          proximal to the hereditary sensory neuropathy type 1 locus on
     RT
     RT
          chromosome 9.";
          Genomics 56:288-295(1999).
     RL
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Application/Control Number: 09/601,582

```
RN
          [6]
          SEQUENCE FROM N.A. (ISOFORM 2A), AND VARIANTS PHE-628 AND ALA-869.
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     RC
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         Martin S.C., Russek S.J., Farb D.H.;
 5
     RA
          "Molecular identification of the human GABABR2: cell surface
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          expression and coupling to adenylyl cyclase in the absence of
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         Mol. Cell. Neurosci. 13:180-191(1999).
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          [7]
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         MEDLINE=99175124; PubMed=10075644;
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         Ng G.Y.K., Clark J., Coulombe N., Ethier N., Hebert T.E., Sullivan R.,
     RA
          Kargman S., Chateauneuf A., Tsukamoto N., McDonald T., Whiting P.,
     RA
          Mezey E., Johnson M.P., Liu Q., Kolakowski L.F. Jr., Evans J.F.,
15
     RA
          Bonner T.I., O'Neill G.P.;
     RA
          "Identification of a GABAB receptor subunit, gb2, required for
     RT
     RT
          functional GABAB receptor activity.";
     RL
          J. Biol. Chem. 274:7607-7610(1999).
20
     RN
          [8]
          R1A-R2 INTERACTION.
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          MEDLINE=20237752; PubMed=10773016;
          Sullivan R., Chateauneuf A., Coulombe N., Kolakowski L.F. Jr.,
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          Johnson M.P., Hebert T.E., Ethier N., Belley M., Metters K.,
     RA
25
          Abramovitz M., O'Neill G.P., Ng G.Y.K.;
     RA
          "Coexpression of full-length gamma-aminobutyric Acid(B) (GABA(B))
     RT
          receptors with truncated receptors and metabotropic glutamate
     RT
          receptor 4 supports the GABA(B) heterodimer as the functional
     RT
     RT
          receptor.";
30
     RL
          J. Pharmacol. Exp. Ther. 293:460-467(2000).
          -!- FUNCTION: RECEPTOR FOR GABA. THE ACTIVITY OF THIS RECEPTOR IS
     CC
              MEDIATED BY G-PROTEINS THAT INHIBITS ADENYLYL CYCLASE ACTIVITY,
     CC
              STIMULATES PHOSPHOLIPASE A2, ACTIVATES POTASSIUM CHANNELS,
     CC
     CC
              INACTIVATES VOLTAGE-DEPENDENT CALCIUM-CHANNELS AND MODULATES
              INOSITOL PHOSPHOLIPIDS HYDROLYSIS. PLAYS A CRITICAL ROLE IN THE
35
     CC
              FINE-TUNING OF INHIBITORY SYNAPTIC TRANSMISSION. PRE-SYNAPTIC
     CC
     CC
              GABA-B-R INHIBIT NEUROTRANSMITTER RELEASE BY DOWN-REGULATING
     CC
              HIGH-VOLTAGE ACTIVATED CALCIUM CHANNELS, WHEREAS POSTSYNAPTIC
              GABA-B-R DECREASE NEURONAL EXCITABILITY BY ACTIVATING A PROMINENT
     CC
40
     CC
              INWARDLY RECTIFYING POTASSIUM (KIR) CONDUCTANCE THAT UNDERLIES THE
     CC
              LATE INHIBITORY POSTSYNAPTIC POTENTIALS. NOT ONLY IMPLICATED IN
     CÇ
              SYNAPTIC INHIBITION BUT ALSO IN HIPPOCAMPAL LONG-TERM
              POTENTIATION, SLOW WAVE SLEEP, MUSCLE RELAXATION AND
     CC
     CC
              ANTINOCICEPTION.
45
          -!- SUBUNIT: HETERODIMER OF GABA-B-R1 AND GABA-B-R2. NEITHER OF WHICH
     CC
              IS EFFECTIVE ON ITS OWN AND HOMODIMERIC ASSEMBLY DOES NOT SEEM TO
     CC
     CC
              HAPPEN.
          -!- SUBCELLULAR LOCATION: INTEGRAL MEMBRANE PROTEIN. MOREOVER
     CC
              COEXPRESSION OF GABA-B-R1 AND GABA-B-R2 APPEARS TO BE A
     CC
50
              PREREQUISITE FOR MATURATION AND TRANSPORT OF GABA-B-R1 TO THE
     CC
     CC
              PLASMA MEMBRANE.
          -!- ALTERNATIVE PRODUCTS: 3 ISOFORMS; 2A (SHOWN HERE), 2B AND 2C; ARE
     CC
     CC
              PRODUCED BY ALTERNATIVE SPLICING.
          -!- TISSUE SPECIFICITY: HIGHLY EXPRESSED IN BRAIN, ESPECIALLY IN
     CC
```

```
CEREBRAL CORTEX, THALAMUS, HIPPOCAMPUS, FRONTAL, OCCIPITAL AND
    CC
             TEMPORAL LOBE, OCCIPITAL POLE AND CEREBELLUM, FOLLOWED BY CORPUS
    CC
             CALLOSUM, CAUDATE NUCLEUS, SPINAL CORD, AMYGDALA AND MEDULLA.
    CC
             WEAKLY EXPRESSED IN HEART, TESTIS AND SKELETAL MUSCLE.
    CC
         -!- DOMAIN: ALPHA-HELICAL PARTS OF THE C-TERMINAL INTRACELLULAR REGION
 5
    CC
            MEDIATE HETERODIMERIC INTERACTION WITH GABA-B RECEPTOR 1.
    CC
          -!- SIMILARITY: BELONGS TO FAMILY 3 OF G-PROTEIN COUPLED RECEPTORS.
    CC
          GABA-B RECEPTOR SUBFAMILY.
    CC
    CC
10
         This SWISS-PROT entry is copyright. It is produced through a
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    collaboration
         between the Swiss Institute of Bioinformatics and the EMBL outstation
         the European Bioinformatics Institute. There are no restrictions on
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     its
         use by non-profit institutions as long as its content is in no
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         modified and this statement is not removed. Usage by and for
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          entities requires a license agreement (See http://www.isb-
    CC
     sib.ch/announce/
         or send an email to license@isb-sib.ch).
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         EMBL; AJ012188; CAA09942.1; -.
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         EMBL; AF056085; AAC63228.1; -.
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         InterPro; IPR000337; GPCR Mgr.
     DR
         Pfam; PF00003; 7tm 3; 1.
     DR
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         Pfam; PF01094; ANF receptor; 1.
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     DR
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     DR
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         PROSITE; PS50099; PRO RICH; 1.
     DR
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     DR
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     DR
          G-protein coupled receptor; Transmembrane; Glycoprotein; Signal;
     KW
          Postsynaptic membrane; Coiled coil; Alternative splicing;
     KW
     KW
          Polymorphism.
50
     FT
          SIGNAL
                       1
                              41
                                       POTENTIAL.
                                       GAMMA-AMINOBUTYRIC ACID TYPE B RECEPTOR,
                      42
                             941
     FT
          CHAIN
                                       SUBUNIT 2.
     FT
     FT
         DOMAIN
                     42
                             483
                                      EXTRACELLULAR (POTENTIAL).
          TRANSMEM 484
                             504
                                     I (POTENTIAL).
     FT
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## Application/Control Number: 09/601,582

	FT	DOMAIN	505	522	CYTOPLASMIC (POTENTIAL).
	FT	TRANSMEM	523	543	II (POTENTIAL).
	FT	DOMAIN	544	551	EXTRACELLULAR (POTENTIAL).
	FT	TRANSMEM	552	572	III POTENTIAL.
5	FT	DOMAIN	573	597	CYTOPLASMIC (POTENTIAL).
	FT	TRANSMEM	598	618	IV (POTENTIAL).
	FT	DOMAIN	619	654	EXTRACELLULAR (POTENTIAL).
	FT	TRANSMEM	655	675	V (POTENTIAL).
	FT	DOMAIN	676	691	CYTOPLASMIC (POTENTIAL).
10	FT	TRANSMEM	692	712	VI (POTENTIAL).
	FT	DOMAIN	713	720	EXTRACELLULAR (POTENTIAL).
	FT	TRANSMEM	721	741	VII (POTENTIAL).
	FT	DOMAIN	742	941	CYTOPLASMIC (POTENTIAL).
	FT	DOMAIN	781	819	COILED COIL (POTENTIAL).
15	FT	CARBOHYD	90	90	N-LINKED (GLCNAC) (POTENTIAL).
	FT	CARBOHYD	298	298	N-LINKED (GLCNAC) (POTENTIAL).
	FT	CARBOHYD	389	389	N-LINKED (GLCNAC) (POTENTIAL).
	FT	CARBOHYD	404	404	N-LINKED (GLCNAC) (POTENTIAL).
	FT	CARBOHYD	453	453	N-LINKED (GLCNAC) (POTENTIAL).
20	FT	VARSPLIC	902	927	MISSING (IN ISOFORM 2B).
	FT	VARSPLIC	929	941	HVPPSFRVMVSGL -> TTLGRGVCCRNTVGSGCGEAGHHG
	FT				WPLRTTRMALRWTGRGRGRLGT (IN ISOFORM 2C).
	FT	VARIANT	628	628	Y -> F.
	FT				/FTId=VAR_010148.
25	FT	VARIANT	869	869	T -> A.
	FT				/FTId=VAR_010149.
	FT	CONFLICT	6	6	$S \rightarrow R (IN REF. 5)$ .
	FT	CONFLICT		12	$P \rightarrow R (IN REF. 5)$ .
	FT	CONFLICT		424	$G \rightarrow E (IN REF. 3).$
30	SQ	SEQUENCE	941 AA;	105821	MW; 09F1773DB0673C5D CRC64;

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08-OCT-1998
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                          5786 bp
            Homo sapiens GABA-B receptor mRNA, complete cds.
DEFINITION
ACCESSION
            AF056085
            g3719225
NID
KEYWORDS
SOURCE
            human.
  ORGANISM
            Homo sapiens
            Eukaryota; Metazoa; Chordata; Vertebrata; Mammalia; Eutheria;
            Primates; Catarrhini; Hominidae; Homo.
               (bases 1 to 5786)
REFERENCE
            Clark, J.C., Lam, A. and Bonner, T.I.
 AUTHORS
            gb2, a second GABA-B receptor
  TITLE
            Unpublished
  JOURNAL
               (bases 1 to 5786)
REFERENCE
            Clark, J.C., Lam, A. and Bonner, T.I.
  AUTHORS
            Direct Submission
  TITLE
            Submitted (27-MAR-1998) Section on Genetics, NIMH, Bldg 36, Rm
  JOURNAL
            3D06, MSC 4094, Bethesda, MD 20892-4094, USA
FEATURES
                     Location/Qualifiers
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                      /chromosome="9"
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                      /tissue type="brain"
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                      /codon start=1
                      /product="GABA-B receptor"
                      /db xref="PID:g3719226"
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                      {\tt GAPRPPPSSPPLSIMGLMPLTKEVAKGSIGRGVLPAVELAIEQIRNESLLRPYFLDLR}
                      \verb|LYDTECDNAKGLKAFYDAIKYGPNHLMVFGGVCPSVTSIIAESLQGWNLVQLSFAATT|
                      \verb"PVLADKKKYPYFFRTVPSDNAVNPAILKLLKHYQWKRVGTLTQDVQRFSEVRNDLTGV"
                      LYGEDIEISDTESFSNDPCTSVKKLKGNDVRIILGQFDQNMAAKVFCCAYEENMYGSK
                      YQWIIPGWYEPSWWEQVHTEANSSRCLRKNLLAAMEGYIGVDFEPLSSKQIKTISGKT
                      PQQYEREYNNKRSGVGPSKFHGYAYDGIWVIAKTLQRAMETLHASSRHQRIQDFNYTD
                      \verb|HTLGRIIL| NAMNETNFFGVTGQVVFRNGERMGTIKFTQFQDSREVKVGEYNAVADTLE|
                      IINDTIRFQGSEPPKDKTIILEQLRKISLPLYSILSALTILGMIMASAFLFFNIKNRN
                      OKLIKMSSPYMNNLIILGGMLSYASIFLFGLDGSFVSEKTFETLCTVRTWILTVGYTT
                      AFGAMFAKTWRVHAIFKNVKMKKKIIKDQKLLVIVGGMLLIDLCILICWQAVDPLRRT
                      VEKYSMEPDPAGRDISIRPLLEHCENTHMTIWLGIVYAYKGLLMLFGCFLAWETRNVS
                      IPALNDSKYIGMSVYNVGIMCIIGAAVSFLTRDQPNVQFCIVALVIIFCSTITLCLVF
                      VPKLITLRTNPDAATQNRRFQFTQNQKKEDSKTSTSVTSVNQASTSRLEGLQSENHRL
                      {\tt RMKITELDKDLEEVTMQLQDTPEKTTYIKQNHYQELNDILNLGNFTESTDGGKAILKN}
                      HLDQNPQLQWNTTEPSRTCKDPIEDINSPEHIQRRLSLQLPILHHAYLPSIGGVDASC
                     VSPCVSPTASPRHRHVPPSFRVMVSGL"
                     5732. .5737
     polyA signal
                     5760. .5765
     polyA signal
                     5786
     polyA site
BASE COUNT
               1415 a
                        1670 c
                                  1474 g
                                           1227 t
ORIGIN
                                   Score 3317.8; DB 12; Length 5786;
  Query Match
                           95.3%;
                                   Pred. No. 0;
                           99.8%;
  Best Local Similarity
                                                                                 1;
  Matches 3332; Conservative
                                  1;
                                      Mismatches
                                                         Indels
                                                                        Gaps
Qу
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Qу	126	cgccgccgcggccgttctgagccgagccggaaccctagcccgagacggagccggggcccg	185
Db		CGCCGCCGCGGCCGTTCTGAGCCGAGCCGGAACCCTAGCCCGAGACGGAGCCGGGGCCCG	
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Db		GGCCGGCGCCATTGCGCGGGGCGCGGGGAAGACCTTGGCGCGGGGCGGGC	410
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Qу			365
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Qу		tgctactgctgctgctgctgctgctctggcgcccggggcctggggctgggcgcggg	425
Db		IGCIACIGCIGCIGCIGCIGCIGCIGCIGCIGCIGCIGCIGCIGC	590 485
Qу			650
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Db		tcgagcagatccgcaacgagtcactcctgcgcccctacttcctcgacctgcggctctatg	605
Qy Db			770
Qу		acacggagtgcgacaacgcaaaagggttgaaagccttctacgatgcaataaaatacgggc	665
Db			830
Qу		cgaaccacttgatggtgtttggaggcgtctgtccatccgtcacatccatc	725
Db			890
Qy	726	ccctccaaggctggaatctggtgcagctttcttttgctgcaaccacgcctgttctagccg	785
Db	891		950
Qy	786	ataaqaaaaatacaccccaccccggaccgcccagacagacaacg	845
Db	951		1010
Qу	846		905
Db	1011		1070
Qу	906	ttcagaggttctctgaggtgcggaatgacctgactggagttctgtatggcgaggacattg	965
Db	1071	TTCAGAGGTTCTCTGAGGTGCGGAATGACCTGACTGGAGTTCTGTATGGCGAGGACATTG	1130
Qу	966	agatttcagacaccgagagcttctccaacgatccctgtaccagtgtcaaaaagctgaagg	1025

.

Db	1131	AGATTTCAGACACCGAGAGCTTCTCCAACGATCCCTGTACCAGTGTCAAAAAGCTGAAGG	1190
Qу	1026	ggaatgatgtgcggatcatccttggccagtttgaccagaatatggcagcaaaagtgttct	1085
Db	1191	GGAATGATGTGCGGATCATCCTTGGCCAGTTTGACCAGAATATGGCAGCAAAAGTGTTCT	1250
Qу	1086	gttgtgcatacgaggagaacatgtatggtagtaaatatcagtggatcattccgggctggt	1145
Db	1251	GTTGTGCATACGAGGAGAACATGTATGGTAGTAAATATCAGTGGATCATTCCGGGCTGGT	1310
Qу	1146	acgagccttcttggtgggagcaggtgcacacggaagccaactcatcccgctgcctccgga	1205
Db	1311	ACGAGCCTTCTTGGTGGGAGCAGGTGCACACGGAAGCCAACTCATCCCGCTGCCTCCGGA	1370
Qу	1206	agaatctgcttgctgccatggagggctacattggcgtggatttcgagcccctgagctcca	1265
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Qу	1266	agcagatcaagaccatctcaggaaagactccacagcagtatgagagaga	1325
Db	1431	AGCAGATCAAGACCATCTCAGGAAAGACTCCACAGCAGTATGAGAGAGA	1490
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Qу	1446	tccaggacttcaactacacggaccacacgctgggcaggatcatcctcaatgccatgaacg	1505
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Qу	1506	agaccaacttcttcggggtcacgggtcaagttgtattccggaatggggagagaatgggga	1565
Db	1671	AGACCAACTTCTTCGGGGTCACGGGTCAAGTTGTATTCCGGAATGGGGAGAATGGGGA	1730
Qу		ccattaaatttactcaatttcaagacagcagggaggtgaaggtgggagagtacaacgctg	1625
Db	1731	CCATTAAATTTACTCAATTTCAAGACAGCAGGGAGGTGAAGGTGGGAGAGTACAACGCTG	1790
Qу		tggccgacacactggagatcatcaatgacaccatcaggttccaaggatccgaaccaccaa	
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Qу		tototgccctcaccatcctcgggatgatcatggccagtgcttttctcttcttcaacatca	
Db	1911	TCTCTGCCCTCACCATCCTCGGGATGATCATGGCCAGTGCTTTTCTCTTCTAACATCA	1970
Qу	1806	agaaccggaatcagaagctcataaagatgtcgagtccatacatgaacaaccttatcatcc	1865
Db		AGAACCGGAATCAGAAGCTCATAAAGATGTCGAGTCCATACATGAACAACCTTATCATCC	
Qу		ttggagggatgctctcctatgcttccatatttctcttttggccttgatggatcctttgtct	
Db	2031	TTGGAGGGATGCTCCCTATGCTTCCATATTTCTCTTTTGGCCTTGATGGATCCTTTGTCT	2090

Qу	1926	ctgaaaagacctttgaaacactttgcaccgtcaggacctggattctcaccgtgggctaca	1985
Db	2091	CTGAAAAGACCTTTGAAACACTTTGCACCGTCAGGACCTGGATTCTCACCGTGGGCTACA	2150
Qу	1986	cgaccgcttttggggccatgtttgcaaagacctggagagtccacgccatcttcaaaaatg	2045
Db	2151	CGACCGCTTTTGGGGGCCATGTTTGCAAAGACCTGGAGAGTCCACGCCATCTTCAAAAATG	2210
Qу		tgaadatgaagaagatgacaccaaggaccagaaaccagagaaccagagagaaagagagagagagagagagagagagagagagag	2105
Db	2211	TGAAAATGAAGAAGAAGATCATCAAGGACCAGAAACTGCTTGTGATCGTGGGGGGCATGC	2270
Qу		[[[]]]	2165
Db		TGCTGATCGACCTGTGTATCCTGATCTGCTGGCAGGCTGTGGACCCCCTGCGAAGGACAG	
Qу		tggagaagtacagcatggagccggacccagcaggacgggatatctccatccgccctctcc	2225
Db		TGGAGAAGTACAGCATGGAGCCGGACCCAGCAGGACGGGATATCTCCATCCGCCCTCTCC	2390
Qу		tggagcactgtgagaacacccatatgaccatctggcttggcatcgtctatgcctacaagg	2285
Db		TGGAGCACTGTGAGAACACCCATATGACCATCTGGCTTGGCATCGTCTATGCCTACAAGG	2450
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Db		GACTTCTCATGTTGTTCGGTTGTTTCTTAGCTTGGGAGACCCGCAACGTCAGCATCCCCG	2510
Qу			2405
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Qу		tcaccctgagaacaaacccagatgcagcaacgcagaacaggcgattccagttcactcaga	
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Qу		acattaaacagaaccactaccaagagctcaatgacatcctcaacctgggaaacttcactg	
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Qу	2826	${\tt agagcacagatggaggaaaggccattttaaaaaaatcacctcgatcaaaatccccagctac}$	2885

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Qу	2946	cagaacacatccagcgtcggctgtccctccagctccccatcctccaccacgcctacctcc	3005
Db	3111		3170
Qу	3006	catccatcggaggcgtggacgccagctgtgtcagcccctgcgtcagccccaccgccagcc	3065
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Qу	3066	cccgccacagacatgtgccaccctccttccgagtcatggtctcgggcctgtaagggtggg	3125
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Qу	3126	gggcctgggcccggggcctcccccgtgacagaaccacactgggcagaggggtctgctgca	3185
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Qy	3186	gaaacactgtcggctctggctgcggagaagctgggcaccatggctggc	3245
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Qу	3366	ttaacatctgcaaacaaggaggcgctgggatatcraatt 3404	
Db	3531		

19-OCT-1998 PRI mRNA 2826 bp LOCUS HSA012188 Homo sapiens mRNA for GABAB receptor, subunit 2. DEFINITION AJ012188 ACCESSION q3776097 NID GABAB receptor; gabab-R2 gene; subunit 2. KEYWORDS SOURCE human. ORGANISM Homo sapiens Eukaryota; Metazoa; Chordata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. (bases 1 to 2826) REFERENCE White, J.H., Wise, A., Main, M.J., Green, A., Fraser, N.J., Disney, G.H., AUTHORS Barnes, A.A., Emson, P., Foord, S.M. and Marshall, F.H. Heterodimerisation is required to form a functional GABAB receptor TITLE Unpublished JOURNAL 2 (bases 1 to 2826) REFERENCE Fraser, N.J. AUTHORS Direct Submission TITLE Submitted (16-OCT-1998) Fraser N.J., Receptor Systems, Cellular **JOURNAL** Sciences, GlaxoWellcome, Medicines Research Centre, Gunnels Wood Road, Stevenage, Herts. SG1 2NY, U.K Location/Qualifiers FEATURES 1. .2826 source /organism="Homo sapiens" /db\_xref="taxon:9606" /tissue type="cerebellum" 1. .114 sig peptide /gene="gabab-R2" 1. .2826 CDS /gene="gabab-R2" /codon start=1 /evidence=experimental /product="GABAB receptor, subunit 2" /db xref="PID:e1333173" /db xref="PID:g3776098" GAPRPPPSSPPLSIMGLMPLTKEVAKGSIGRGVLPAVELAIEQIRNESLLRPYFLDLR LYDTECDNAKGLKAFYDAIKYGPNHLMVFGGVCPSVTSIIAESLQGWNLVQLSFAATT PVLADKKKYPYFFRTVPSDNAVNPAILKLLKHYQWKRVGTLTQDVQRFSEVRNDLTGV LYGEDIEISDTESFSNDPCTSVKKLKGNDVRIILGQFDQNMAAKVFCCAYEENMYGSK YQWIIPGWYEPSWWEQVHTEANSSRCLRKNLLAAMEGYIGVDFEPLSSKQIKTISGKT PQQYEREYNNKRSGVGPSKFHGYAYDGIWVIAKTLQRAMETLHASSRHQRIQDFNYTD HTLGRIILNAMNETNFFGVTGQVVFRNGERMGTIKFTQFQDSREVKVGEYNAVADTLE IINDTIRFQGSEPPKDKTIILEQLRKISLPLYSILSALTILGMIMASAFLFFNIKNRN QKLIKMSSPYMNNLIILGGMLSYASIFLFGLDGSFVSEKTFETLCTVRTWILTVGYTT AFGAMFAKTWRVHAIFKNVKMKKKIIKDQKLLVIVGGMLLIDLCILICWQAVDPLRRT VEKYSMEPDPAGRDISIRPLLEHCENTHMTIWLGIVYAYKGLLMLFGCFLAWETRNVS IPALNDSKYIGMSVYNVGIMCIIGAAVSFLTRDQPNVQFCIVALVIIFCSTITLCLVF VPKLITLRTNPDAATQNRRFQFTQNQKKEDSKTSTSVTSVNQASTSRLEGLQSENHRL RMKITELDKDLEEVTMQLQDTPEKTTYIKQNHYQELNDILNLGNFTESTDGGKAILKN HLDONPOLOWNTTEPSRTCKDPIEDINSPEHIQRRLSLQLPILHHAYLPSIGGVDASC VSPCVSPTASPRHRHVPPSFRVMVSGL" 1. .2826 gene /gene="gabab-R2" mat\_peptide 115. .2823 /gene="gabab-R2" /product="GABAB receptor, subunit 2" 697 a 841 c 719 g 569 t BASE COUNT ORIGIN

100.0%; Pred. No. 0; Best Local Similarity Gaps 0; 0; Mismatches 1; Indels Matches 2825; Conservative Qy Db 353 gcgcgcctgctactgctactgctgctgctgctgctgctctggcgcccggggcctgg 412 Qу 61 GCGCGCCTGCTACTGCTGCTGCTGCCGCTGCTGCCTCTGGCGCCCCGGGGCCTGG 120 Db 413 ggctgggcgcgggggcgcccccggccgccagcagccgccgctctccatcatgggc 472 Qу 121 GGCTGGGCGGGGGCGCCCCCGGCCGCCCAGCAGCCCGCCGCTCTCCATCATGGGC 180 Db 473 ctcatqccqctcaccaaggagqtggccaagggcagcatcgggcgcggtgtgctccccgcc 532 Qу 181 CTCATGCCGCTCACCAAGGAGGTGGCCAAGGGCAGCATCGGGCGCGGTGTGCTCCCCGCC 240 Db 533 qtggaactggccatcgagcagatccgcaacgagtcactcctgcgcccctacttcctcgac 592 Qу 241 GTGGAACTGGCCATCGAGCAGATCCGCAACGAGTCACTCCTGCGCCCCTACTTCCTCGAC 300 Db 593 ctgcggctctatgacacggagtgcgacaacgcaaaagggttgaaagccttctacgatgca 652 QУ 301 CTGCGGCTCTATGACACGGAGTGCGACAACGCAAAAGGGTTGAAAGCCTTCTACGATGCA 360 Db 653 ataaaatacgggccgaaccacttgatggtgtttggaggcgtctgtccatccgtcacatcc 712 0ν 361 ATAAAATACGGGCCTAACCACTTGATGGTGTTTTGGAGGCGTCTGTCCATCCGTCACATCC 420 Db Qy Db 773 cctgttctagccgataagaaaaatacccttatttctttcggaccgtcccatcagacaat 832 Qу 481 CCTGTTCTAGCCGATAAGAAAAATACCCTTATTTCTTTCGGACCGTCCCATCAGACAAT 540 Db 833 gcggtgaatccagccattctgaagttgctcaagcactaccagtggaagcgcgtgggcacg 892 Qу 541 GCGGTGAATCCAGCCATTCTGAAGTTGCTCAAGCACTACCAGTGGAAGCGCGTGGGCACG 600 Db 893 ctgacgcaagacgttcagaggttctctgaggtgcggaatgacctgactggagttctgtat 952 Ov 601 CTGACGCAAGACGTTCAGAGGTTCTCTGAGGTGCGGAATGACCTGACTGGAGTTCTGTAT 660 Db 953 ggcgaggacattgagatttcagacaccgagagcttctccaacgatccctgtaccagtgtc 1012 Qу 661 GGCGAGGACATTGAGATTTCAGACACCGAGAGCTTCTCCAACGATCCCTGTACCAGTGTC 720 Db 1013 aaaaagctgaaggggaatgatgtgcggatcatccttggccagtttgaccagaatatggca 1072 Qу 721 AAAAAGCTGAAGGGGAATGATGTGCGGATCATCCTTGGCCAGTTTGACCAGAATATGGCA 780 Db 1073 gcaaaagtqttctqttqtqcatacqagqagaacatgtatggtagtaaatatcagtggatc 1132 Qу 781 GCAAAAGTGTTCTGTTGTGCATACGAGGAGAACATGTATGGTAGTAAATATCAGTGGATC 840 Db 1133 attccgggctggtacgagccttcttggtgggagcaggtgcacacggaagccaactcatcc 1192 Qy 

Db	841	ATTCCGGGCTGGTACGAGCCTTCTTGGTGGGAGCCAGGTGCACACGGAAGCCAACTCATCC	900
Qу	1193	cgctgcctccggaagaatctgcttgctgccatggagggctacattggcgtggatttcgag	1252
Db	901		960
Qy	1253	cccctgagctccaagcagatcaagaccatctcaggaaagactccacagcagtatgagaga	1312
Db	961		1020
Qy	1313	Uau Lacadedacadedacedacedagedagedagedagedagedagedagedagedagedag	1372
Db	1021		1080
Qу	1373	ggcatctgggtcatcgccaagacactgcagagggccatggagacactgcatgccagcagc	1432
Db	1081		1140
Qу	1433	cggcaccagcggatccaggacttcaactacacggaccacacgctgggcaggatcatcctc	1492
Db	1141		1200
Qу	1493	aatgccatgaacgagaccaacttcttcggggtcacgggtcaagttgtattccggaatggg	1552
Db	1201	AATGCCATGAACGAGACCAACTTCTTCGGGGTCACGGGTCAAGTTGTATTCCGGAATGGG	1260
Qу	1553	gagagaatggggaccattaaatttactcaatttcaagacagcagggaggtgaaggtggga	1612
Db	1261	GAGAGAATGGGGACCATTAAATTTACTCAATTTCAAGACAGCAGGGAGGTGAAGGTGGGA	1320
Qу	1613	gagtacaacgctgtggccgacacactggagatcatcaatgacaccatcaggttccaagga	1672
Db	1321	GAGTACAACGCTGTGGCCGACACACTGGAGATCATCAATGACACCATCAGGTTCCAAGGA	1380
Qу	1673	tccgaaccaccaaaagacaagaccatcatcctggagcagctgcggaagatctccctacct	1732
Db	1381	TCCGAACCACCAAAAGACAAGACCATCATCCTGGAGCAGCTGCGGAAGATCTCCCTACCT	1440
Qу	1733	ctctacagcatcctctctgccctcaccatcctcgggatgatcatggccagtgcttttctc	1792
Db	1441	CTCTACAGCATCCTCTGCCCTCACCATCCTCGGGATGATCATGGCCAGTGCTTTTCTC	1500
Qу	1793	ttcttcaacatcaagaaccggaatcagaagctcataaagatgtcgagtccatacatgaac	1852
Db	1501	TTCTTCAACATCAAGAACCGGAATCAGAAGCTCATAAAGATGTCGAGTCCATACATGAAC	1560
Qу	1853	aaccttatcatccttggagggatgctctcctatgcttccatatttctcttttggccttgat	1912
Db	1561	AACCTTATCATCCTTGGAGGGATGCTCTCCTATGCTTCCATATTTCTCTTTGGCCTTGAT	1620
Qу	1913	ggatcctttgtctctgaaaagacctttgaaacactttgcaccgtcaggacctggattctc	1972
Db	1621	GGATCCTTTGTCTCTGAAAAGACCTTTGAAACACTTTGCACCGTCAGGACCTGGATTCTC	1680
Qу	1973	accgtgggctacacgaccgcttttggggccatgtttgcaaagacctggagagtccacgcc	2032
Db	1681	ACCGTGGGCTACACGACCGCTTTTGGGGCCATGTTTGCAAAGACCTGGAGAGTCCACGCC	1740
Qу		atcttcaaaaatgtgaaaatgaagaagaagatcatcaaggaccagaaactgcttgtgatc	
Db	1741	ATCTTCAAAAATGTGAAAATGAAGAAGAAGATCATCAAGGACCAGAAACTGCTTGTGATC	1800

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Qу	2093	gtggggggcatgctgctgatcgacctgtgtatcctgatctgctggcaggctgtggacccc	2152
Db	1801	GTGGGGGGCATGCTGATCGACCTGTGTATCCTGATCTGCTGGCAGGCTGTGGACCCC	1860
Qу	2153	ctgcgaaggacagtggagaagtacagcatggagccggacccagcaggacgggatatctcc	2212
Db	1861	CTGCGAAGGACAGTGGAGAAGTACAGCATGGAGCCGGACCCAGCAGGACGGATATCTCC	1920
Qу	2213	atccgccctctcctggagcactgtgagaacacccatatgaccatctggcttggcatcgtc	2272
Db	1921	ATCCGCCCTCTCCTGGAGCACTGTGAGAACACCCATATGACCATCTGGCTTGGCATCGTC	1980
Qу	2273	tatgcctacaagggacttctcatgttgttcggttgtttcttagcttgggagacccgcaac	2332
Db	1981	TATGCCTACAAGGGACTTCTCATGTTGTTCGGTTGTTTCTTAGCTTGGGAGACCCGCAAC	2040
Qу	2333	gtcagcatccccgcactcaacgacagcaagtacatcgggatgagtgtctacaacgtgggg	2392
Db	2041	GTCAGCATCCCCGCACTCAACGACAGCAAGTACATCGGGATGAGTGTCTACAACGTGGGG	2100
Qу	2393	atcatgtgcatcatcggggccgctgtctccttcctgacccgggaccagcccaatgtgcag	2452
Db	2101	ATCATGTGCATCATCGGGGCCGCTGTCTCCTTCCTGACCCGGGACCAGCCCAATGTGCAG	2160
Qу	2453	ttctgcatcgtggctctggtcatcatcttctgcagcaccatcaccctctgcctggtattc	2512
Db	2161	TTCTGCATCGTGGCTCTGGTCATCATCTTCTGCAGCACCATCACCCTCTGCCTGGTATTC	2220
Qу	2513	gtgccgaagctcatcaccctgagaacaaacccagatgcagcaacgcagaacaggcgattc	2572
Db	2221	GTGCCGAAGCTCATCACCCTGAGAACAAACCCAGATGCAGCAACGCAGAACAGGCGATTC	2280
Qу	2573	cagttcactcagaatcagaagaagaagattctaaaacgtccacctcggtcaccagtgtg	2632
Db	2281	CAGTTCACTCAGAATCAGAAGAAGAAGAATTCTAAAACGTCCACCTCGGTCACCAGTGTG	2340
Qу	2633	aaccaagccagcacatcccgcctggagggcctacagtcagaaaaccatcgcctgcgaatg	2692
Db	2341	AACCAAGCCAGCACATCCCGCCTGGAGGGCCTACAGTCAGAAAACCATCGCCTGCGAATG	2400
Qу	2693	aagatcacagagctggataaagacttggaagaggtcaccatgcagctgcaggacacaca	2752
Db	2401	AAGATCACAGAGCTGGATAAAGACTTGGAAGAGGTCACCATGCAGCTGCAGGACACACCA	2460
Qу	2753	gaaaagaccacctacattaaacagaaccactaccaagagctcaatgacatcctcaacctg	2812
Db	2461	GAAAAGACCACCTACATTAAACAGAACCACTACCAAGAGCTCAATGACATCCTCAACCTG	2520
Qу	2813	ggaaacttcactgagagcacagatggaggaaaggccattttaaaaaaatcacctcgatcaa	2872
Db	2521	GGAAACTTCACTGAGAGCACAGATGGAGGAAAGGCCATTTTAAAAAAATCACCTCGATCAA	2580
Qу	2873	aatccccagctacagtggaacacaacagagccctctcgaacatgcaaagatcctatagaa	2932
Db	2581	AATCCCCAGCTACAGTGGAACACAGAGCCCTCTCGAACATGCAAAGATCCTATAGAA	2640
Qу	2933	gatataaactctccagaacacatccagcgtcggctgtccctccagctccccatcctccac	2992
Db	2641	GATATAAACTCTCCAGAACACATCCAGCGTCGGCTGTCCCTCCAGCTCCCCATCCTCCAC	2700
Qу	2993	${\tt cacgcctacctcccatccatcggaggcgtggacgccagctgtgtcagcccctgcgtcagc}$	3052

Db	2701		2760
Qу		cccaccgccagccccgccacagacatgtgccaccctccttccgagtcatggtctcgggc	
Db	2761	CCCACCGCCAGCCCCGCCACAGACATGTGCCACCCTCCTTCCGAGTCATGGTCTCGGGC	2820
Qу	3113	ctgtaa 3118 	
Db	2821	CTGTAA 2826	

15-OCT-1998 5459 bp mRNA ROD LOCUS AF058795 Rattus norvegicus GABA-B receptor gb2 mRNA, complete cds. DEFINITION ACCESSION AF058795 g3746525 NID KEYWORDS Norway rat. SOURCE ORGANISM Rattus norvegicus Eukaryota; Metazoa; Chordata; Vertebrata; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus. (bases 1 to 5459) REFERENCE AUTHORS Clark, J.C., Lam, A. and Bonner, T.I. gb2, A second GABA-B receptor TITLE Unpublished JOURNAL (bases 1 to 5459) REFERENCE Clark, J.C., Lam, A. and Bonner, T.I. AUTHORS TITLE Direct Submission Submitted (09-APR-1998) Section on Genetics, NIMH, Bldg. 36, Rm. **JOURNAL** 3D06, MSC 4094, Bethesda, MD 20892-4094, USA **FEATURES** Location/Qualifiers 1. .5459 source /organism="Rattus norvegicus" /db xref="taxon:10116" /tissue type="cerebral cortex" 151. .2973 CDS /note="G protein-coupled receptor; similar to Rattus norvegicus GABA-B receptor GABA-BR1b encoded by Genbank Accession Number Y10370; similar to Homo sapiens gb2 GABA-B receptor encoded by Genbank Accession Number AF056085" /codon start=1 /product="GABA-B receptor gb2" /db xref="PID:g3746526" /translation="MASPPSSGQPRPPPPPPPPPPPARLLLPLLLSLLLWLAPGAWGWTRG  $\verb|APRPPPSSPPLSIMGLMPLTKEVAKGSIGRGVLPAVELAIEQIRNESLLRPYFLDLRL|$ YDTECDNAKGLKAFYDAIKYGPNHLMVFGGVCPSVTSIIAESLQGWNLVQLSFAATTP VLADKKKYPYFFRTVPSDNAVNPAILKLLKHFRWRRVGTLTQDXQRFSEVRNDLTGVL YGEDIEISDTESFSNDPCTSVKKLKGNDVRIILGQFDQNMAAKVFCCAFEESMFGSKY QWIIPGWYEPAWWEQVHVEANSSRCLRRSLLAAMEGYIGVDFEPLSSKQIKTISGKTP QQFEREYNSKRSGVGPSKFHGYAYDGIWVIAKTLQRAMETLHASSRHQRIQDFNYTDH TLGKIILNAMNETNFFGVTGQVVFRNGERMGTIKFTQFQDSREVKVGEYNAVADTLEI INDTIRFQGSEPPKDKTIILEQLRKISLPLYSILSALTILGMIMASAFLFFNIKNRNQ KLIKMSSPYMNNLIILGGMLSYASIFLFGLDGSFVSEKTFETLCTVRTWILTVGYTTA FGAMFAKTWRVHAIFKNVKMKKKIIKDQKLLVIVGGMLLIDLCILICWQAVDPLRRTV ERYSMEPDPAGRDISIRPLLEHCENTHMTIWLGIVYAYKGLLMLFGCFLAWETRNVSI PALNDSKYIGMSVYNVGIMCIIGAAVSFLTRDQPNVQFCIVALVIIFCSTITLCLVFV PKLITLRTNPDAATQNRRFQFTQNQKKEDSKTSTSVTSVNQASTSRLEGLQSENHRLR MKITELDKDLEEVTMQLQDTPEKTTYIKQNHYQELNDILSLGNFTESTDGGKAILKNH LDONPOLOWNTTEPSRTCKDPIEDINSPEHIQRRLSLQLPILHHAYLPSIGGVDASCV SPCVSPTASPRHRHVPPSFRVMVSGL" 5421. .5426 polyA signal polyA\_site 5459 BASE COUNT 1334 a 1603 c 1395 g 1120 t 7 others ORIGIN 72.9%; Score 2536.6; DB 13; Length 5459; Query Match Best Local Similarity 88.1%; Pred. No. 0; 1; Mismatches 355; Indels 30; Gaps 7; Matches 2849; Conservative

Qy 165 ccgagacggagccggggcccgggcgcattgcgcgggcgccgcgggaagaccttgg 224

Db	23	CCGGAGCCGAGCCGGGCCCGTGCCGGCGCCATTGCGCGGGGCGCCGCGGGCAAAKCTCGG	82
Qу		cgcggggcgggccgggccaggccatgcgggccgagtgagccggcgccgcagcccgc	
Db	83	CGCCGGGCGGGCCAGGCCATGCGGGCCGAGTGAGCTGGCGCCCGCAGCCCGC	142
Qу		ggcgcggcatggcttccccgcggagctccgggcagcccgggccgccgccgccgccac	
Db	143	GGCGCGGCATGGCTTCCCCGCCGAGCTCCGGGCAGCCCCGGCCGCCGCCGCCGCCGCCGCCGCCGCC	199
Qу		cgccgcccgcgcctgctactgctactgctgctgccgctgctgctctggcgcccg	
Db	200	CGCCGCCGCGCGCGCTGCTGCTGCTGCTGCTGCTGCTGCT	259
Qу		gggcctggggctgggcgcgcgcccccggccgccagcagccgccgctctcca	
Db	260	GGGCCTGGGGCTGGACGCGGGGGGGCGCCCCCGGCCGCCCAGCAGCCCGCCG	319
Qу	465	tcatgggcctcatgccgctcaccaaggaggtggccaagggcagcatcgggcgcggtgtgc	524
Db	320	TCATGGGCCTCATGCCGCTCACCAAGGAGGTGGCCAAGGGCAGCATCGGGCGCGTGC	379
Qу	525	tccccgccgtggaactggccatcgagcagatccgcaacgagtcactcctgcgcccctact	584
Db	380	TCCCCGCCGTGGAGCTAGCCATCGAGCAGATCCGCAACGAGTCACTCCTGCGCCCCTACT	439
Qу	585	tcctcgacctgcggctctatgacacggagtgcgacaacgcaaaagggttgaaagccttct	644
Db	440	TCCTGGACCTGCGACTCTATGACACCGAGTGTGACAATGCAAAGGGACTGAAAGCCTTCT	499
Qу	645	acgatgcaataaaatacgggccgaaccacttgatggtgtttggaggcgtctgtccatccg	704
Db	500	ATGACGCAATAAAGTATGGGCCGAACCATTTGATGGTGTTTTGGAGGCGTCTGTCCGTCTG	559
Qу	705	tcacatccatcattgcagagtccctccaaggctggaatctggtgcagctttcttt	764
Db	560	TCACATCTATTATCGCGGAGTCCCTCCAAGGCTGGAATCTGGTGCAGCTTTCCTTCGCCG	619
Qу	765	caaccacgcctgttctagccgataagaaaaatacccttatttctttc	824
Db	620	CCACCACGCCTGTTCTTGCGGATAAGAAGAAGTACCCGTATTTCTTCCGGACGGTGCCGT	679
Qу	825	cagacaatgcggtgaatccagccattctgaagttgctcaagcactaccagtggaagcgcg	884
Db	680	CAGACAACGCGGTGAACCCCGCCATCCTGAAGCTCCTGAAGCACTTCCGCTGGCGGCGTG	739
Qу	885	tgggcacgctgacgcaagacgttcagaggttctctgaggtgcggaatgacctgactggag	944
Db	740	TGGGCACACTCACGCAGGACGYGCAGCGCTTCTCCGAGGTGAGGAATGACCTGACTGGGG	799
Qу	945	ttctgtatggcgaggacattgagatttcagacaccgagagcttctccaacgatccctgta	1004
Db	800	TTCTGTATGGGGAAGATATTGAGATCTCAGACACAGAGAGTTTCTCCAATGATCCCTGCA	859
Qу	1005	ccagtgtcaaaaagctgaaggggaatgatgtgcggatcatccttggccagtttgaccaga	1064
Db	860	CCAGCGTCAAAAAGCTCAAGGGGAATGACGTGCGGATCATCCTTGGCCAGTTTGACCAGA	919
Qy	1065	atatggcagcaaaagtgttctgttgtgcatacgaggagaacatgtatggtagtaaatatc	1124
Dh	920	ATATGGCAGCAAAAGTCTTCTGTTGTGCCTTCGAGGAGAGCATGTTTGGCAGCAAGTACC	979

Qу		agtggatcattccgggctggtacgagccttcttggtgggagcaggtgcacacggaagcca	
Db	980	AGTGGATCATCCCGGGATGGTACGAGCCTGCGTGGTGGAGCAGGTGCATGTGGAGGCCA	1039
Qу		actcatcccgctgcctccggaagaatctgcttgctgccatggagggctacattggcgtgg	
Db		ATTCCTCACGCTGCCTGCGCAGAAGCCTCCTGGCTGCCATGGAAGGTTACATCGGAGTGG	
Qу		atttcgagccctgagctccaagcagatcaagaccatctcaggaaagactccacagcagt	
Db		ACTTTGAGCCCCTGAGCTCCAAACAAATCAAGACCATCTCAGGGAAGACTCCACAGCAGT	
Qу		atgagagagtacaacaacaagcggtcaggcgtggggcccagcaagttccacgggtacg	
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Qу		cctacgatggcatctgggtcatcgccaagacactgcagagggccatggagacactgcatg	
Db		CCTACGATGGGATCTGGGTCATCGCCAAGACCCTACAGAGGGCCATGGAGACACTGCATG	
Qу		ccagcagccggcaccagcggatccaggacttcaactacacggaccacacgctgggcagga	
Db		CCAGTAGCAGGCACCAGCGGATCCAGGACTTCAACTACACAGACCACACGCTGGGCAAAA	
Qу		tcatcctcaatgccatgaacgagaccaacttcttcggggtcacgggtcaagttgtattcc	
Db		TCATCCTCAATGCCATGAACGAGACCAACTTCTTCGGGGGTCACGGGTCAAGTTGTGTTCCC	
Qу		ggaatggggagaatggggaccattaaatttactcaatttcaagacagcagggaggtga	
Db		GGAACGGGGAGAATGGGAACCATTAAATTTACTCAATTTCAAGACAGCAGAGAGGTGA	
Qу		aggtgggagagtacaacgctgtggccgacacactggagatcatcaatgacaccatcaggt	
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Qу		tccaaggatccgaaccaccaaaagacaagaccatcatcctggagcagctgcggaagatct	
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Qу		ccctacctctctacagcatcctctctgccctcaccatcctcgggatgatcatggccagtg	
Db		CGCTTCCACTGTATAGCATCCTGTCCGCTCTCACCATCCTCGGCATGATCATGGCCAGCG	
Qу		cttttctcttcttcaacatcaagaaccggaatcagaagctcataaagatgtcgagtccat	
Db		CCTTCCTCTTCAACATCAAGAACCGGAACCAAAAGCTGATTAAGATGTCAAGCCCCT	
Qу		acatgaacaaccttatcatccttggagggatgctctcctatgcttccatatttctctttg	
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Qу		gccttgatggatcctttgtctctgaaaagacctttgaaacactttgcaccgtcaggacct	
Db		GCCTCGACGGGTCCTTCGTCTCAGAAAAGACCTTTGAAACACTCTGCACGGTCCGGACCT	
Qу		ggattctcaccgtgggctacacgaccgcttttggggccatgtttgcaaagacctggagag	
Db		GGATTCTCACCGTGGGCTACACAACTGCCTTTGGGGCCATGTTTGCAAAGACCTGGAGGG	
Ov	2025	tccacgccatcttcaaaaatgtgaaaatgaagaagaagatcatcaaggaccagaaactgc	. 2004

Db	1880		1939
Qу	2085	ttgtgatcgtggggggcatgctgctgatcgacctgtgtatcctgatctgctggcaggctg	2144
Db	1940	TTGTGATTGTGGGGGGCATGCTCATCGACCTGTGCATCCTGATCTGTTGGCAGGCTG	1999
Qу	2145	tggacccctgcgaaggacagtggagaagtacagcatggagccggacccagcaggacggg	2204
Db	2000	TGGACCCCTGCGGAGGACAGTGGAGAGGTACAGCATGGAGCCGGACCCAGCAGGCCGGG	2059
Qу	2205	atatctccatccgccctctcctggagcactgtgagaacacccatatgaccatctggcttg	2264
Db	2060	ACATCTCCATCCGCCCATTGCTGGAACACTGCGAAAACACCCACATGACCATCTGGCTTG	2119
Qу	2265	gcatcgtctatgcctacaagggacttctcatgttgttcggttgtttcttagcttgggaga	2324
Db	2120	GCATTGTCTACGCCTACAAGGGGCTCCTCATGCTATTCGGTTGTTTCTTGGCATGGGAAA	2179
Qу	2325	cccgcaacgtcagcatccccgcactcaacgacagcaagtacatcgggatgagtgtctaca	2384
Db	2180	CCCGCAATGTGAGCATCCCTGCCCTCAACGACAGCAAGTACATCGGCATGAGTGTGTACA	2239
Qу	2385	acgtggggatcatgtgcatcatcggggccgctgtctccttcct	2444
Db	2240	ATGTGGGGATCATGTGCATCATCGGGGCTGCTGTCTCCTTCCT	2299
Qy		<pre>atgtgcagttctgcatcgtggctctggtcatcatcttctgcagcaccatcaccctctgcc                                      </pre>	
Db		ACGTGCAGTTCTGCATCGTGGCCCTGGTCATCTTCTGCAGCACCATCACTCTCTGCC	
Qу		tggtattcgtgccgaagctcatcaccctgagaacaaacccagatgcagcaacgcagaaca	
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08-OCT-1998 PRI mRNA 743 bp LOCUS AF095723 Homo sapiens GABA-B receptor splice variant 1 mRNA, partial cds. DEFINITION AF095723 ACCESSION q3719471 NTD KEYWORDS SOURCE human. ORGANISM Homo sapiens Eukaryota; Metazoa; Chordata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 743) REFERENCE Clark, J.C., Lam, A. and Bonner, T.I. AUTHORS TITLE gb2, a second GABA-B receptor Unpublished JOURNAL 2 (bases 1 to 743) REFERENCE Clark, J.C., Lam, A. and Bonner, T.I. AUTHORS Direct Submission TITLE Submitted (27-MAR-1998) Section on Genetics, National Institute of JOURNAL Mental Health, Bldg 36, Rm 3D06, MSC 4094, Bethesda, MD 20892-4090, USA Location/Qualifiers **FEATURES** 1. .743 source /organism="Homo sapiens" /db xref="taxon:9606" /tissue\_type="brain" /map = "D9S287 - D9S176"/chromosome="9" <1. .549 CDS /note="alternative splice variant of GABA-B receptor encoded by GenBank Accession Number AF056085; G protein-coupled receptor; gb2" /codon start=1 /product="GABA-B receptor splice variant 1" /db xref="PID:g3719472" /translation="ITLCLVFVPKLITLRTNPDAATQNRRFQFTQNQKKEDSKTSTSV TSVNOASTSRLEGLOSENHRLRMKITELDKDLEEVTMQLQDTPEKTTYIKQNHYQELN DILNLGNFTESTDGGKAILKNHLDQNPQLQWNTTEPSRTCKDPIEDINSPEHIQRRLS LQLPILHHRHVPPSFRVMVSGL" 225 c 181 g 123 t 214 a BASE COUNT ORIGIN Score 653.4; DB 12; Length 743; 18.8%; Query Match Best Local Similarity 90.4%; Pred. No. 1.3e-95; 78; 1; Indels Gaps Matches 742; Conservative 0; Mismatches 1; 2492 atcaccctctgcctggtattcgtgccgaagctcatcaccctgagaacaaacccagatgca 2551 Qу 1 ATCACCCTCTGCCTGGTATTCGTGCCGAAGCTCATCACCCTGAGAACAAACCCAGATGCA 60

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GenBank (R) AF069755 LOCUS (LOC): GenBank ACC. NO. (GBN): AF069755 222290-10-0 CAS REGISTRY NO. (RN): SEQUENCE LENGTH (SQL): 3075 mRNA; linear MOLECULE TYPE (CI): Primates DIVISION CODE (CI): 7 Apr 1999 DATE (DATE): Homo sapiens orphan G protein-coupled receptor DEFINITION (DEF): HG20 (HG20) mRNA, complete cds. human. SOURCE: Homo sapiens ORGANISM (ORGN): Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo NUCLEIC ACID COUNT (NA): 729 a 925 c 830 q 591 t 1 (bases 1 to 3075) REFERENCE: Ng, G.Y.K.; McDonald, T.; Bonnert, T.; Rigby, M.; AUTHOR (AU): Heavens, R.; Whiting, P.; Chateauneuf, A.; Coulombe, N.; Kargmen,S.; Caskey,T.; Evans,J.; O'Neill,G.P.; Liu,Q. Cloning of a novel G-protein-coupled receptor GPR 51 TITLE (TI): resembling GABAB receptors expressed predominantly in nervous tissues and mapped proximal to the hereditary sensory neuropathy type 1 locus on chromosome 9 Genomics, 56 (3), 288-295 (1999) JOURNAL (SO): 2 (bases 1 to 3075) REFERENCE: McDonald, T.; Liu, Q. AUTHOR (AU): Direct Submission TITLE (TI): Submitted (03-JUN-1998) Human Genetics, Merck Research JOURNAL (SO): Labs, West Point, PA 19486, USA FEATURES (FEAT): Qualifier Location Feature Key \_\_\_\_\_\_+ /organism="Homo sapiens" 1..3075 source /db-xref="taxon:9606" /chromosome="9" /map="4.81 cM from WI-8684" /gene="HG20" 1..3075 gene CDS 157..2982 /gene="HG20" /note="member of GABAb receptor family" /codon-start=1 /product="orphan G protein-coupled receptor HG20" /protein-id="AAC99345.1" /db-xref="PID:g4091933" /db-xref="GI:4091933" /translation="MASPRRSGQPGRPPPPPPPP ARLLLLLLPLLPLAPGAWGWAR GAPRPPPSSPPLSIMGLMPLTKEVAKGSIGRGVL PAVELAIEQIRNESLLRPYFLDLR LYDTECDNAKGLKAFYDAIKYGPNHLMVFGGVCP SVTSIIAESLQGWNLVQLSFAATT PVLADKKKYPYFFRTVPSDNAVNPAILKLLKHYQ WKRVGTLTQDVQRFSEVRNDLTGV LYGEDIEISDTESFSNDPCTSVKKLKGNDVRIIL GOFDONMAAKVFCCAYEENMYGSK

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ACCESSION AB016161 NID g3929914

KEYWORDS rat GABAB receptor 1d.

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ORGANISM Rattus norvegicus

Eukaryota; Metazoa; Chordata; Vertebrata; Mammalia; Eutheria;

Rodentia; Sciurognathi; Muridae; Murinae; Rattus.

REFERENCE 1 (sites)

AUTHORS Isomoto, S., Kaibara, M., Yano, K. and Taniyama, K.

TITLE Cloning and tissue distribution of novel isoform of rat GABAB

receptors

JOURNAL Biochem. Biophys. Res. Commun. (1998) In press

REFERENCE 2 (bases 1 to 3202)

AUTHORS Isomoto, S.

TITLE Direct Submission

JOURNAL Submitted (10-JUL-1998) to the DDBJ/EMBL/GenBank databases. Shojiro

Isomoto, Nagasaki University School of Medicine, Third Department of Internal Medicine; Sakamoto 1-7-1, Nagasaki, Nagasaki 852-8102,

Japan (Tel:+81-95-849-7047, Fax:+81-95-849-7048)

FEATURES Location/Qualifiers

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ORIGIN

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        121 IKYGPNHLMVFGGVCPSVTSIIAESLQGWNLVQLSFAATTPVLADKKKYPYFFRTVPSDN 180
Db
     181 AVNPAILKLLKHYQWKRVGTLTQDVQRFSEVRNDLTGVLYGEDIEISDTESFSNDPCTSV 240
Qу
         181 AVNPAILKLLKHYQWKRVGTLTQDVQRFSEVRNDLTGVLYGEDIEISDTESFSNDPCTSV 240
Db
     241 KKLKGNDVRIILGQFDQNMAAKVFCCAYEENMYGSKYQWIIPGWYEPSWWEQVHTEANSS 300
Qу
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Db
     301 RCLRKNLLAAMEGYIGVDFEPLSSKQIKTISGKTPQQYEREYNNKRSGVGPSKFHGYAYD 360
Qу
         301 RCLRKNLLAAMEGYIGVDFEPLSSKQIKTISGKTPQQYEREYNNKRSGVGPSKFHGYAYD 360
Db
     361 GIWVIAKTLQRAMETLHASSRHQRIQDFNYTDHTLGRIILNAMNETNFFGVTGQVVFRNG 420
Qу
         361 GIWVIAKTLQRAMETLHASSRHQRIQDFNYTDHTLGRIILNAMNETNFFGVTGQVVFRNG 420
Db
     421 ERMGTIKFTQFQDSREVKVGEYNAVADTLEIINDTIRFQGSEPPKDKTIILEQLRKISLP 480
Qу
         421 ERMGTIKFTQFQDSREVKVGEYNAVADTLEIINDTIRFQGSEPPKDKTIILEQLRKISLP 480
Db
     481 LYSILSALTILGMIMASAFLFFNIKNRNQKLIKMSSPYMNNLIILGGMLSYASIFLFGLD 540
Qу
         481 LYSILSALTILGMIMASAFLFFNIKNRNQKLIKMSSPYMNNLIILGGMLSYASIFLFGLD 540
Db
     541 GSFVSEKTFETLCTVRTWILTVGYTTAFGAMFAKTWRVHAIFKNVKMKKKIIKDQKLLVI 600
Qу
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Qу	3082	gccaccctccttccgagtcatggtctcgggcctgtaagggtgggggcctgggcccgggg	3141
Db	326	GCCACCCTCCTTCCGANTCATGGTCTCGGGCCTGTAANGGTGGGGGGCCTGGGCCCGGGG	267
Qу	3142	cctccccgtgacagaaccacactgggcagaggggtctgctgcagaaacactgtcggctc	3201
Db	266	CCTCCCCCGTGACAGAACCACACTGGGCAGAGGGGTCTGCTGCAGAAANACTGTCGGCTC	207
Qу	3202	tggctgcggagaagctgggcaccatggctggcctctcaggaccactcggatggcactcag	3261
Db	206	TGGCTGCGGAGAANCTGGGCACCATGGCTGGCCTCTCAGGACCACTCGGATGGCACTCAG	147
Qу	3262	gtggacaggacggggcagggggagacttggcacctgacctcgagccttatttgtgaagtc	3321
Db	146	GTGGACAGGACGGGGCAGGGGGAGACTTGGCACCTGACCTCGAGCCTTATTTGTGAAGTC	87
Qу	3322	cttatttcttcacaaagaaggaacggaaatgggacgtcttccttaacatctgcaaaca	3381
Db	86	CTTATTCTCACAAAGAAGAAGAACGGAAATGGGACGTCTTCCTTAACATCTGCAAACA	27
Qу	3382	aggaggcgctgggatatcraatt 3404	
Db	26		

Db	541	GSFVSEKTFETLCTVRTWILTVGYTTAFGAMFAKTWRVHAIFKNVKMKKKIIKDQKLLVI	600
Qу	601	VGGMLLIDLCILICWQAVDPLRRTVEKYSMEPDPAGRDISIRPLLEHCENTHMTIWLGIV	660
Db	601	VGGMLLIDLCILICWQAVDPLRRTVEKYSMEPDPAGRDISIRPLLEHCENTHMTIWLGIV	660
Qу	-	YAYKGLLMLFGCFLAWETRNVSIPALNDSKYIGMSVYNVGIMCIIGAAVSFLTRDQPNVQ	
Db	661	YAYKGLLMLFGCFLAWETRNVSIPALNDSKYIGMSVYNVGIMCIIGAAVSFLTRDQPNVQ	720
Qу		FCIVALVIIFCSTITLCLVFVPKLITLRTNPDAATQNRRFQFTQNQKKEDSKTSTSVTSV	
Db	721	FCIVALVIIFCSTITLCLVFVPKLITLRTNPDAATQNRRFQFTQNQKKEDSKTSTSVTSV	780
Qу		NQASTSRLEGLQSENHRLRMKITELDKDLEEVTMQLQDTPEKTTYIKQNHYQELNDILNL	
Db	781	NQASTSRLEGLQSENHRLRMKITELDKDLEEVTMQLQDTPEKTTYIKQNHYQELNDILNL	840
Qу		GNFTESTDGGKAILKNHLDQNPQLQWNTTEPSRTCKDPIEDINSPEHIQRRLSLQLPILH	
Db	841	GNFTESTDGGKAILKNHLDQNPQLQWNTTEPSRTCKDPIEDINSPEHIQRRLSLQLPILH	900
Qу		HAYLPSIGGVDASCVSPCVSPTASPRHRHVPPSFRVMVSGL 941	
Db	901	HAYLPSIGGVDASCVSPCVSPTASPRHRHVPPSFRVMVSGL 941	

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PRT:
                                  940 AA.
   088871
             PRELIMINARY;
ID
AC
    088871;
   01-NOV-1998 (TREMBLREL. 08, CREATED)
01-NOV-1998 (TREMBLREL. 08, LAST SEQUENCE UPDATE)
01-NOV-1998 (TREMBLREL. 08, LAST ANNOTATION UPDATE)
DT
DT
DT
    GABA-B RECEPTOR GB2.
DE
    RATTUS NORVEGICUS (RAT).
OS
    EUKARYOTA; METAZOA; CHORDATA; VERTEBRATA; MAMMALIA; EUTHERIA;
OC
    RODENTIA; SCIUROGNATHI; MURIDAE; MURINAE; RATTUS.
OC
RN
    SEQUENCE FROM N.A.
RP
    TISSUE=CEREBRAL CORTEX;
RC
    CLARK J.C., LAM A., BONNER T.I.;
RA
    "gb2, A second GABA-B receptor.";
RT
    SUBMITTED (APR-1998) TO EMBL/GENBANK/DDBJ DATA BANKS.
RL
    EMBL; AF058795; G3746526; -.
DR
            940 AA; 105762 MW; 94C83CC1 CRC32;
    SEOUENCE
SO
                     97.9%; Score 4422.5; DB 10; Length 940;
 Query Match
 Best Local Similarity 97.6%; Pred. No. 0;
                                                               1;
                                                     1; Gaps
 Matches 918; Conservative 16; Mismatches
                                          6;
                                             Indels
      1 MASPRSSGQPGPPPPPPPPPPARLLLLLLLLLLLLPLLPLAPGAWGWARGAPRPPPSSPPLSIMG 60
Qу
        1 MASPPSSGQP-RPPPPPPPPPPARLLLPLLLSLLLWLAPGAWGWTRGAPRPPPSSPPLSIMG 59
Dh
      61 LMPLTKEVAKGSIGRGVLPAVELAIEQIRNESLLRPYFLDLRLYDTECDNAKGLKAFYDA 120
Qу
        60 LMPLTKEVAKGSIGRGVLPAVELAIEQIRNESLLRPYFLDLRLYDTECDNAKGLKAFYDA 119
Db
     121 IKYGPNHLMVFGGVCPSVTSIIAESLQGWNLVQLSFAATTPVLADKKKYPYFFRTVPSDN 180
Qу
        120 IKYGPNHLMVFGGVCPSVTSIIAESLQGWNLVQLSFAATTPVLADKKKYPYFFRTVPSDN 179
Db
     181 AVNPAILKLLKHYQWKRVGTLTQDVQRFSEVRNDLTGVLYGEDIEISDTESFSNDPCTSV 240
Qу
        180 AVNPAILKLLKHFRWRRVGTLTQDXQRFSEVRNDLTGVLYGEDIEISDTESFSNDPCTSV 239
Db
     241 KKLKGNDVRIILGQFDQNMAAKVFCCAYEENMYGSKYQWIIPGWYEPSWWEQVHTEANSS 300
Qу
        240 KKLKGNDVRIILGQFDQNMAAKVFCCAFEESMFGSKYQWIIPGWYEPAWWEQVHVEANSS 299
Db
     301 RCLRKNLLAAMEGYIGVDFEPLSSKQIKTISGKTPQQYEREYNNKRSGVGPSKFHGYAYD 360
Qу
        300 RCLRRSLLAAMEGYIGVDFEPLSSKQIKTISGKTPQQFEREYNSKRSGVGPSKFHGYAYD 359
Db
     361 GIWVIAKTLQRAMETLHASSRHQRIQDFNYTDHTLGRIILNAMNETNFFGVTGQVVFRNG 420
Qу
        360 GIWVIAKTLQRAMETLHASSRHQRIQDFNYTDHTLGKIILNAMNETNFFGVTGQVVFRNG 419
Db
     421 ERMGTIKFTQFQDSREVKVGEYNAVADTLEIINDTIRFQGSEPPKDKTIILEQLRKISLP 480
Qy
         420 ERMGTIKFTQFQDSREVKVGEYNAVADTLEIINDTIRFQGSEPPKDKTIILEQLRKISLP 479
Db
     481 LYSILSALTILGMIMASAFLFFNIKNRNQKLIKMSSPYMNNLIILGGMLSYASIFLFGLD 540
Qу
         480 LYSILSALTILGMIMASAFLFFNIKNRNQKLIKMSSPYMNNLIILGGMLSYASIFLFGLD 539
Db
     541 GSFVSEKTFETLCTVRTWILTVGYTTAFGAMFAKTWRVHAIFKNVKMKKKIIKDQKLLVI 600
Qу
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Db	540	GSFVSEKTFETLCTVRTWILTVGYTTAFGAMFAKTWRVHAIFKNVKMKKKIIKDQKLLVI	599
Qу	601	VGGMLLIDLCILICWQAVDPLRRTVEKYSMEPDPAGRDISIRPLLEHCENTHMTIWLGIV	660
Db	600	VGGMLLIDLCILICWQAVDPLRRTVERYSMEPDPAGRDISIRPLLEHCENTHMTIWLGIV	659
Qу	661	YAYKGLLMLFGCFLAWETRNVSIPALNDSKYIGMSVYNVGIMCIIGAAVSFLTRDQPNVQ	720
Db	660	YAYKGLLMLFGCFLAWETRNVSIPALNDSKYIGMSVYNVGIMCIIGAAVSFLTRDQPNVQ	719
QУ		FCIVALVIIFCSTITLCLVFVPKLITLRTNPDAATQNRRFQFTQNQKKEDSKTSTSVTSV	
Db	720	FCIVALVIIFCSTITLCLVFVPKLITLRTNPDAATQNRRFQFTQNQKKEDSKTSTSVTSV	779
QУ	781	NQASTSRLEGLQSENHRLRMKITELDKDLEEVTMQLQDTPEKTTYIKQNHYQELNDILNL	
Db	780	THE THE PARTY AND THE PROPERTY OF THE PROPERTY	839
Qу		GNFTESTDGGKAILKNHLDQNPQLQWNTTEPSRTCKDPIEDINSPEHIQRRLSLQLPILH	
Db	840	GNFTESTDGGKAILKNHLDQNPQLQWNTTEPSRTCKDPIEDINSPEHIQRRLSLQLPILH	899
Qy		HAYLPSIGGVDASCVSPCVSPTASPRHRHVPPSFRVMVSGL 941	
Db	900	HAYLPSIGGVDASCVSPCVSPTASPRHRHVPPSFRVMVSGL 940	

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PRT;
                                     844 AA.
               PRELIMINARY;
    008621
АC
    008621;
    01-JUL-1997 (TREMBLREL. 04, CREATED)
DT
    01-JUL-1997 (TREMBLREL. 04, LAST SEQUENCE UPDATE)
DΤ
    01-NOV-1998 (TREMBLREL. 08, LAST ANNOTATION UPDATE)
DΤ
    GABA-BR1B RECEPTOR.
DE
GN
    GABA-BR1B.
    RATTUS NORVEGICUS (RAT).
OS
    EUKARYOTA; METAZOA; CHORDATA; VERTEBRATA; MAMMALIA; EUTHERIA; RODENTIA;
OC
    SCIUROGNATHI; MURIDAE; MURINAE; RATTUS.
OC
RN
    SEQUENCE FROM N.A.
RΡ
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RC
    MEDLINE; 97222131.
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    KAUPMANN K., HEGGEL K., HEID J., FLOR P.J., BISCHOFF S., MICKEL S.J.,
    MCMASTER G., ANGST C., BITTIGER H., FROESTL W., BETTLER B.;
    "Expression cloning of GABA(B) receptors uncovers similarity to
    metabotropic glutamate receptors.";
    NATURE 386:239-246(1997).
    EMBL; Y10370; E311196; -.
    PFAM; PF01094; ANF receptor; 1.
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                       25.4%; Score 1146; DB 10; Length 844;
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 Matches 294; Conservative 188; Mismatches 294; Indels
                                                        54; Gaps
                                                                     18;
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Qу
        3 PGGPCTPVGWP-----LPLLLVMAAGVAPVWASHSPHLPRPHPRVPPHPSSERRAVY 54
Db
      58 IMGLMPLTKEVAKGSIGRGVLPAVELAIEQIRN-ESLLRPYFLDLRLYDTECDNAKGLKA 116
Qy
         55 IGALFPMS---GGWPGGQACQPAVEMALEDVNSRRDILPDYELKLIHHDSKCDPGQATKY 111
Db
     117 FYDAIKYGPNHLMVFGGVCPSVTSIIAESLQGWNLVQLSFAATTPVLADKKKYPYFFRTV 176
Qу
         :|: : | ::: | |:||:||:::||: : | ||:||::::| |::: ::| |||||
     112 LYELLYNDPIKIILMPG-CSSVSTLVAEAARMWNLIVLSYGSSSPALSNRQRFPTFFRTH 170
Db
     177 PSDNAVNPAILKLLKHYQWKRVGTLTQDVQRFSEVRNDLTGVLYGEDIEISDTESFSNDP 236
Qу
             ||::||: ||:::|:| : |: :|| :
                                                  171 PSATLHNPTRVKLFEKWGWKKIATIQQTTEVFTSTLDDLEERVKEAGIEITFRQSFFSDP 230
Db
     237 CTSVKKLKGNDVRIILGQFDQNMAAKVFCCAYEENMYGSKYQWIIPGWYEPSWWEQVHTE 296
QУ
          :||:|| | |||:|| : | |||| || ||::|||| ||:||
     231 AVPVKNLKRQDARIIVGLFYETEARKVFCEVYKERLFGKKYVWFLIGWYADNWF---KTY 287
Db
     297 ANSSRCLRKNLLAAMEGYIGVDFEPLSSKQIKTISGKTPQQYEREYNN--KRSGVGPSKF 354
Qу
          | | :: |:|| | : |:: ::|| |:|::
                                                : 11
     288 DPSINCTVEEMTEAVEGHITTEIVMLNPANTRSISNMTSQEFVEKLTKRLKRHPEETGGF 347
Db
     355 H--GYAYDGIWVIAKTLQRAMETLHASSRH-QRIQDFNYTDHTLGRIILNAMNETNFFGV 411
Qу
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Db
     412 TGQVVF-RNGERMGTIKFTQFQDSREVKVGEYNAVADTLEIINDTIRFQGSEPPKDKTII 470
Qу
         :|:||| :| ||:
                        405 SGHVVFDASGSRMAWTLIEQLQGGSYKKIGYYDSTKDDLS-WSKTDKWIGGSPPADQTLV 463
Db
     471 LEOLRKISLPLYSILSALTILGMIMASAFLFFNIKNRNQKLIKMSSPYMNNLIILGGMLS 530
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Qу

Db	464	: : :   : :   :   :::         ::     :   :   :    :  :	523
Qу	531	YASIFLFGLDGSFVSEKTFETLCTVRTWILTVGYTTAFGAMFAKTWRVHAIFKNVKMKK-	590
Db	524	LAAVFPLGLDGYHIGRSQFPFVCQARLWLLGLGFSLGYGSMFTKIWWVHTVFTKKEEKKE	583
Qу	590	KIIKDQKLLVIVGGMLLIDLCILICWQAVDPLRRTVEKYSMEPDPAGRDISIRPLLEH	647
Db	584	WRKTLEPWKLYATVGLLVGMDVLTLAIWQIVDPLHRTIETFAKEEPKEDIDVSILPQLEH	643
Qу	648	CENTHMTIWLGIVYAYKGLLMLFGCFLAWETRNVSIPALNDSKYIGMSVYNVGIMCIIGA	707
Db	644	CSSKKMNTWLGIFYGYKGLLLLLGIFLAYETKSVSTEKINDHRAVGMAIYNVAVLCLITA	703
Qу	708	AVSFLTRDQPNVQFCIVALVIIFCSTITLCLVFVPKLITLRTNPDAATQNRRFQFTQNQK : :	767
Db	704	PVTMILSSQQDAAFAFASLAIVFSSYITLVVLFVPKMRRLITRGEWQSET	753
Qy	768	KEDSKTSTSVTSVNQASTSRLEGLQSENHRLRMKITELDKDLEEVTMQLQ 817 :   ::   :	
Db	754	QDTMKTGSS-TNNNEEEKSRLLEKENRELEKIIAEKEERVSELRHQLQ 800	

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V10265 standard; cDNA to mRNA; 2620 BP.
ΙD
AC
    03-JUN-1998 (first entry)
DT
    Human GABA-BR1a/b receptor cDNA.
DE
    Gamma-animobutyric acid; GABA-BRla/b receptor; human; brain; agonist;
KW
    inhibitory neurotransmitter; peripheral nervous system; antagonist;
KW
    treatment; dementia; depression; anxiety; bronchial inflammation; asthma;
KW
    epilepsy; cognitive function; ds.
KW
OS
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                  Location/Qualifiers
FH
    Key
                  1. .2382
FT
    CDS
FT
                  /*tag= a
                  /product= GABA-bR1a/b
FT
PN
    WO9746675-A1.
    11-DEC-1997.
PD
    19-MAR-1997; E01370.
ΡF
PR
    22-NOV-1996; US-756091.
PR
    30-MAY-1996; US-655716.
    (NOVS ) NOVARTIS AG.
PΑ
    Bettler B, Bittiger H, Froestl W, Kaupmann K, Mickel SJ;
PΙ
    WPI; 98-042183/04.
    P-PSDB; W40117.
    Purified GABA-B receptor or receptor protein - and antagonists of
    these which may be useful in treating nervous system disorders
PΤ
    Claim 3; Page 56-62; 108pp; English. → №.79.86
PS
    This cDNA sequence encodes a novel human GABA-B receptor protein,
CC
    GABA-BR1a/b. GABA (gamma-aminobutyric acid) is the major inhibitory
CC
    neurotransmitter found in the brain and peripheral nervous system
CC
    and this receptor may be used for the identification of GABA-B
CC
    receptor agonists and antagonists. Such proteins may be used in
CC
    treatment of dementia, depression, anxiety, epilepsy, spasticity,
CC
    bronchial inflammation or asthma or to improve cognitive function.
CC
    GABA-B receptor ligands and probes derived from this sequence can be
CC
    used to assay for GABA-B receptors or DNA encoding them.
CC
                                 716 C;
                                          710 G;
SQ
    Sequence
             2620 BP;
                        593 A;
                       9.0%; Score 313.6; DB 1; Length 2620;
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                             0; Mismatches 1004; Indels
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Qу
      64 CCCGCGGTGGAGATGGCGCTGGAGGACGTGAATAGCCGCAGGGACATCCTGCCGGACTAT 123
Db
     584 tteetegaeetgeggetetatgaeaeggagtgegaeaaegeaaagggttgaaageette 643
Qу
                     124 GAGCTCAAGCTCATCCACCACGACAGCAAGTGTGATCCAGGCCAAGCCACCAAGTACCTA 183
Db
     644 tacgatgcaataaaatacgggccgaaccacttgatggtgttttggaggcgtctgtccatcc 703
Qу
                 184 TATGAGCTGCTCTACAACGACCCTATCAA---GATCATCCTTATGCCTGGCTGCAGCTCT 240
     Qу
                                     241 GTCTCCACGCTGGTGGCTGAGGCTGCTAGGATGTGGAACCTCATTGTGCTTTCCTATGGC 300
Db
     Qу
         301 TCCAGCTCACCAGCCCTGTCAAACCGGCAGCGTTTCCCCACTTTCTTCCGAACGCACCCA 360
Db
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Qу		tcagacaatgcggtgaatccagccattctgaagttgctcaagcactaccagtggaagcgc	
Db		TCAGCCACACTCCACAACCCTACCCGCGTGAAACTCTTTGAAAAGTGGGGCTGGAAGAAG	
Qу		gtgggcacgctgacgcaagacgttcagaggttctctgaggtgcggaatgacctgactgga	
Db	421	ATTGCTACCATCCAGCAGACCACTGAGGTCTTCACTTCGACTCTGGACGACCTGGAGGAA	480
Qу	944	gttctgtatggcgaggacattgagatttcagacaccgagagcttctccaacgatccctgt	1003
Db	481	CGAGTGAAGGAGGCTGGAATTGAGATTACTTTCCGCCAGAGTTTCTTCTCAGATCCAGCT	540
Qу	1004	accagtgtcaaaaagctgaaggggaatgatgtgcggatcatccttggccagtttgaccag	1063
Db	541	GTGCCCGTCAAAAACCTGAAGCGCCAGGATGCCCGAATCATCGTGGGACTTTTCTATGAG	600
Qу	1064	aatatggcagcaaaagtgttctgttgtgcatacgaggagaacatgtatggtagtaaatat	1123
Db	601	ACTGAAGCCCGGAAAGTTTTTTGTGAGGTGTACAAGGAGCGTCTCTTTGGGAAGAAGTAC	660
Qу	1124	cagtggatcattccgggctggtacgagccttcttggtgggagcaggtgcacacggaagcc	1183
Db	661	GTCTGGTTCCTCATTGGGTGGTATGCTGACAATTGGTTCAAGATCTACGAC	711
Qу	1184	<pre>aactcatcccgctgcctccggaagaatctgcttgctgccatggagggctacattggcgtg</pre>	1243
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Qу	1244	gatttcgagcccctgagctccaagcagatcaagaccatctcaggaaagactccacagcag	1303
Db	772	GAGATTGTCATGCTGAATCCTGCCAATACCCGCAGCATTTCCAACATGACATCCCAGGAA	831
Qу	1304	tatgagagagagtacaacaacaagcggtcaggcgtggggcccagcaagttccac	1357
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Qу	1358	<pre>gggtacgcctacgatggcatctgggtcatcgccaagacactgcagagggccatggagaca  </pre>	1417
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Qу	1418	ctgcatgccagcagccggcaccagcggatccaggacttcaactacacggaccacacgctg	1477
Db	952	GGAGGAGGCGGCCGTTCTGGTGTGCGCCTGGAGGACTTCAACTACAACAACCAGACCATT	1011
Qу	1478	ggcaggatcatcctcaatgccatgaacgagaccaacttcttcggggtcacgggtcaagtt	1537
Db	1012	ACCGACCAAATCTACCGGGCAATGAACTCTTCGTCCTTTGAGGGTGTCTCTGGCCATGTG	1071
Qy	1538	gtattccggaatggggagagaatggggaccattaaatttactcaatttcaagacagg	1597
Db	1072		1131
Qу	1598	gaggtgaaggtgggagagtacaacgctgtggccgacacactggagatcatcaatgacacc	1657
Db	1132		1191
Qу	1658	atcaggttccaaggatccgaaccaccaaaagacaagaccatcatcctggagcagctgcgg	1717
Db	1192	GATAAATGGATTGGAGGGTCCCCCCAGCTGACCAGACCCTGGTCATCAAGACATTCCGC	1251
Qv	1718	aagatctccctacctctacagcatcctctctgccctcaccatcctcgggatgatcatg	1777

Db	1252		1311
Qу	1778	gccagtgcttttctcttcttcaacatcaagaaccggaatcagaagctcataaagatgtcg	1837
Db	1312	GCTGTTGTCTGTCCTTTAACATCTACAACTCACATGTCCGTTATATCCAGAACTCA	1371
Qу	1838	agtccatacatgaacaaccttatcatccttggagggatgctctcctatgcttccatattt	1897
Db	1372	CAGCCCAACCTGAACAACCTGACTGCTGTGGGCTGCTCACTGGCTTTAGCTGCTGTCTTC	1431
Qу	1898	ctctttggccttgatggatcctttgtctctgaaaagacctttgaaacactttgcaccgtc	1957
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Qу	1958	aggacctggattctcaccgtgggctacacgaccgcttttggggccatgtttgcaaagacc	2017
Db	1492	CGCCTCTGGCTCCTGGGCTTTAGTCTGGGCTACGGTTCCATGTTCACCAAGATT	1551
Qу	2018	tggagagtccacgccatcttcaaaaatgtgaaaatgaagaagaagatcatcaaggacc	2076
Db	1552	TGGTGGGTCCACACGGTCTTCACAAAGAAGAAGAAGAAGAAGAGAGTGGAGGAAGAACTCTG	1611
Qу	2076	agaaactgcttgtgatcgtggggggcatgctgctgatcgacctgtgtatcctg	2128
Db	1612	GAACCCTGGAAGCTGTATGCCACAGTGGGCCTGCTGGTGGGCATGGATGTCCTCACTCTC	1671
Qу	2129	atctgctggcaggctgtggaccccctgcgaaggacagtggagaagtacagcatggagccg	2188
Db	1672	GCCATCTGGCAGATCGTGGACCCTCTGCACCGGACCATTGAGACATTTGCCAAGGAGAA	1731
Qу	2189	gacccagcaggacgggatatctccatccgccctctcctggagcactgtgagaacacccat	2248
Db	1732	CCTAAGGAAGATATTGACGTCTCTATTCTGCCCCAGCTGGAGCATTGCAGCTCCAGGAAG	1791
Qу	2249	atgaccatctggcttggcatcgtctatgcctacaagggacttctcatgttgttcggttgt	2308
Db	1792	ATGAATACATGGCTTGGCATTTTCTATGGTTACAAGGGGCTGCTGCTGCTGCTGGGAATC	1851
Qу	2309	ttcttagcttgggagacccgcaacgtcagcatccccgcactcaacgacagcaagtacatc	2368
Db	1852	TTCCTTGCTTATGAGACCAAGAGTGTGTCCACTGAGAAGATCAATGATCACCGGGCTGTG	1911
Qу	2369	gggatgagtgtctacaacgtggggatcatgtgcatcatcggggccgctgtctccttcct	2428
Db	1912	GGCATGCTATCTACAATGTGGCAGTCCTGTGCCTCATCACTGCTCCTGTCACCATGATT	1971
Qу	2429	acccgggaccagcccaatgtgcagttctgcatcgtggctctggtcatcatcttctgcagc	2488
Db	1972		2031
Qу	2489	accatcaccctctgcctggtattcgtgccgaagctcatcaccctgagaac 2538	
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W40118 standard; Protein; 844 AA.
ΙD
AC
    W40118;
    03-JUN-1998 (first entry)
\mathsf{DT}
    Rat GABA-BR1b receptor protein.
DE
    Gamma-animobutyric acid; GABA-BR1b receptor; rat; brain; agonist;
KW
    inhibitory neurotransmitter; peripheral nervous system; antagonist;
KW
    treatment; dementia; depression; anxiety; bronchial inflammation; asthma;
KW
    epilepsy; cognitive function.
KW
    Rattus norvegicus.
OS
    WO9746675-A1.
PN
    11-DEC-1997.
PD
    19-MAR-1997; E01370.
ΡF
    22-NOV-1996; US-756091.
PR
    30-MAY-1996; US-655716.
PR
PΑ
    (NOVS ) NOVARTIS AG.
    Bettler B, Bittiger H, Froestl W, Kaupmann K, Mickel SJ;
ΡI
    WPI; 98-042183/04.
DR
    N-PSDB; V10266.
DR
    Purified GABA-B receptor or receptor protein - and antagonists of
PΤ
    these which may be useful in treating nervous system disorders
PT
    Claim 4; Page 74-79; 108pp; English.
PS
    This sequence represents a novel rat GABA-B receptor protein,
CC
    GABA-BR1b. GABA (gamma-aminobutyric acid) is the major inhibitory
CC
    neurotransmitter found in the brain and peripheral nervous system
CC
    and this receptor may be used for the identification of GABA-B
CC
    receptor agonists and antagonists. Such proteins may be used in
CC
    treatment of dementia, depression, anxiety, epilepsy, spasticity,
CC
    bronchial inflammation or asthma or to improve cognitive function.
CC
    GABA-B receptor ligands and probes derived from this sequence can be
    used to assay for GABA-B receptors or DNA encoding them.
CC
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    Sequence
                        25.4%; Score 1146; DB 1; Length 844;
  Query Match
  Best Local Similarity 35.4%; Pred. No. 1.9e-109;
                                                                      18;
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QУ
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Db
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Db
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         :|: : | ::: | |:||::::||: : |||: ||:::::| |::: ::| ||||
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     237 CTSVKKLKGNDVRIILGQFDQNMAAKVFCCAYEENMYGSKYQWIIPGWYEPSWWEQVHTE 296
Qу
           231 AVPVKNLKRQDARIIVGLFYETEARKVFCEVYKERLFGKKYVWFLIGWYADNWF---KTY 287
Db
      297 ANSSRCLRKNLLAAMEGYIGVDFEPLSSKQIKTISGKTPQQYEREYNN--KRSGVGPSKF 354
Qу
           | | :: |:|| | : |:: ::|| |:|:: : ||
      288 DPSINCTVEEMTEAVEGHITTEIVMLNPANTRSISNMTSQEFVEKLTKRLKRHPEETGGF 347
Db
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Qу		HGYAYDGIWVIAKTLQRAMETLHASSRH-QRIQDFNYTDHTLGRIILNAMNETNFFGV :    :  : :  :  :   ::	
Db		QEAPLAYDAIWALALALNKTSGGGGRSGVRLEDFNYNNQTITDQ1YRAMNSSSFEGV	
Qу		TGQVVF-RNGERMGTIKFTQFQDSREVKVGEYNAVADTLEIINDTIRFQGSEPPKDKTII : :   :   :   :   :   :   :   :   :   :	
Db	405	SGHVVFDASGSRMAWTLIEQLQGGSYKKIGYYDSTKDDLS-WSKTDKWIGGSPPADQTLV	463
Qу		LEQLRKISLPLYSILSALTILGMIMASAFLFFNIKNRNQKLIKMSSPYMNNLIILGGMLS : :  :    :   :     :   :     :   :     :	
Db	464	IKTFRFLSQKLFISVSVLSSLGIVLAVVCLSFNIYNSHVRYIQNSQPNLNNLTAVGCSLA	523
QУ	531	YASIFLFGLDGSFVSEKTFETLCTVRTWILTVGYTTAFGAMFAKTWRVHAIFKNVKMKK-	590
Db	524	LAAVFPLGLDGYHIGRSQFPFVCQARLWLLGLGFSLGYGSMFTKIWWVHTVFTKKEEKKE	583
Qу	590	KIIKDQKLLVIVGGMLLIDLCILICWQAVDPLRRTVEKYSMEPDPAGRDISIRPLLEH	647
Db	584	WRKTLEPWKLYATVGLLVGMDVLTLAIWQIVDPLHRTIETFAKEEPKEDIDVSILPQLEH	643
Qy	648	CENTHMTIWLGIVYAYKGLLMLFGCFLAWETRNVSIPALNDSKYIGMSVYNVGIMCIIGA	707
Db	644	CSSKKMNTWLGIFYGYKGLLLLLGIFLAYETKSVSTEKINDHRAVGMAIYNVAVLCLITA	703
Qу	708	AVSFLTRDQPNVQFCIVALVIIFCSTITLCLVFVPKLITLRTNPDAATQNRRFQFTQNQK : : :   :   :   !           : :     :   :	767
Db	704	: : :   :   :   :       ::    :     ::  PVTMILSSQQDAAFAFASLAIVFSSYITLVVLFVPKMRRLITRGE	753
Qу	768	KEDSKTSTSVTSVNQASTSRLEGLQSENHRLRMKITELDKDLEEVTMQLQ 817	
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V10267 standard; cDNA to mRNA; 2924 BP.
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AC
    03-JUN-1998 (first entry)
DΤ
    Human GABA-BR1b receptor cDNA.
DF.
    Gamma-animobutyric acid; GABA-BRla/b receptor; human; brain; agonist;
KW
    inhibitory neurotransmitter; peripheral nervous system; antagonist;
KW
    treatment; dementia; depression; anxiety; bronchial inflammation; asthma;
ΚW
    epilepsy; cognitive function; ds.
KW
OS
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                  Location/Qualifiers
FH
    Key
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FT
    CDS
                  /*tag= a
FT
                  /product= GABA-BR1b
FT
    WO9746675-A1.
PN
PD
    11-DEC-1997.
    19-MAR-1997; E01370.
PF
    22-NOV-1996; US-756091.
PR
    30-MAY-1996; US-655716.
PR
    (NOVS ) NOVARTIS AG.
PΑ
    Bettler B, Bittiger H, Froestl W, Kaupmann K, Mickel SJ;
PΙ
    WPI; 98-042183/04.
DR
    P-PSDB; W40119.
DR
    Purified GABA-B receptor or receptor protein - and antagonists of
PΤ
    these which may be useful in treating nervous system disorders
PT
    Claim 3; Page 79-86; 108pp; English.
PS
    This cDNA sequence encodes a novel human GABA-B receptor protein,
CC
    GABA-BR1b. GABA (gamma-aminobutyric acid) is the major inhibitory
CC
    neurotransmitter found in the brain and peripheral nervous system
СC
    and this receptor may be used for the identification of GABA-B
CC
    receptor agonists and antagonists. Such proteins may be used in
CC
    treatment of dementia, depression, anxiety, epilepsy, spasticity,
CC
    bronchial inflammation or asthma or to improve cognitive function.
CC
    GABA-B receptor ligands and probes derived from this sequence can be
CC
    used to assay for GABA-B receptors or DNA encoding them.
CC
                                         793 G;
            2924 BP;
                        628 A;
                                852 C;
SO
    Sequence
                       9.0%; Score 313.6; DB 1; Length 2924;
  Query Match
 Best Local Similarity 49.1%; Pred. No. 6.9e-48;
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                            0; Mismatches 1004; Indels
                                                       30; Gaps
  Matches 996; Conservative
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Qу
         385 CCCGCGGTGGAGATGGCGCTGGAGGACGTGAATAGCCGCAGGGACATCCTGCCGGACTAT 444
Db
     584 ttcctcgacctgcggctctatgacacggagtgcgacaacgcaaaagggttgaaagccttc 643
Qу
                    1.1
           445 GAGCTCAAGCTCATCCACCACGACAGCAAGTGTGATCCAGGCCAAGCCACCAAGTACCTA 504
Db
     644 tacgatgcaataaaatacgggccgaaccacttgatggtgtttggaggcgtctgtccatcc 703
Qу
               11
     505 TATGAGCTGCTCTACAACGACCCTATCAA---GATCATCCTTATGCCTGGCTGCAGCTCT 561
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     Qу
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Db
     Qу
          622 TCCAGCTCACCAGCCCTGTCAAACCGGCAGCGTTTCCCCACTTTCTTCCGAACGCACCCA 681
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Qу	824	tcagacaatgcggtgaatccagccattctgaagttgctcaagcactaccagtggaagcgc	883
Db	682	TCAGCCACACCCCACAACCCTACCCGCGTGAAACTCTTTGAAAAGTGGGGCTGGAAGAAG	741
Qу	884	gtgggcacgctgacgcaagacgttcagaggttctctgaggtgcggaatgacctgactgga	943
Db	742	ATTGCTACCATCCAGCAGACCACTGAGGTCTTCACTTCGACTCTGGACGACCTGGAGGAA	801
Qу	944	gttctgtatggcgaggacattgagatttcagacaccgagagcttctccaacgatccctgt	1003
Db	802	CGAGTGAAGGAGGCTGGAATTGAGATTACTTTCCGCCAGAGTTTCTTCTCAGATCCAGCT	861
Qу	1004	accagtgtcaaaaagctgaaggggaatgatgtgcggatcatccttggccagtttgaccag	1063
Db	862	GTGCCCGTCAAAAACCTGAAGCGCCAGGATGCCCGAATCATCGTGGGACTTTTCTATGAG	921
Qу	1064	aatatggcagcaaaagtgttctgttgtgcatacgaggagaacatgtatggtagtaaatat	1123
Db	922	ACTGAAGCCCGGAAAGTTTTTTGTGAGGTGTACAAGGAGCGTCTCTTTGGGAAGAAGTAC	981
Qу	1124	<pre>cagtggatcattccgggctggtacgagccttcttggtgggagcaggtgcacacggaagcc</pre>	1183
Db	982	GTCTGGTTCCTCATTGGGTGGTATGCTGACAATTGGTTCAAGATCTACGAC	1032
Qу	1184	<pre>aactcatcccgctgcctccggaagaatctgcttgctgccatggagggctacattggcgtg</pre>	1243
Db	1033	CCTTCTATCAACTGCACAGTGGATGAGATGACTGAGGCGGTGGAGGGCCACATCACAACT	1092
Qу	1244	<pre>gatttcgagcccctgagctccaagcagatcaagaccatctcaggaaagactccacagcag                                       </pre>	1303
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Db	1153	TTTGTGGAGAAACTAACCAAGCGACTGAAAAGACACCCTGAGGAGACAGGAGGCTTCCAG	1212
Qу	1358	gggtacgcctacgatggcatctgggtcatcgccaagacactgcagagggccatggagaca	1417
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Qу	1418	ctgcatgccagcagcagcaccagcggatccaggacttcaactacacggaccacacgctg	1477
Db	1273	GGAGGAGGCGGCCGTTCTGGTGTGCGCCTGGAGGACTTCAACTACAACAACCAGACCATT	1332
Qу	1478	ggcaggatcatcctcaatgccatgaacgagaccaacttcttcggggtcacgggtcaagtt	1537
Db	1333	ACCGACCAAATCTACCGGGCAATGAACTCTTCGTCCTTTGAGGGTGTCTCTGGCCATGTG	1392
Qy	1538	gtattccggaatggggagaatggggaccattaaatttactcaatttcaagacagcagg	1597
Db	1393	GTGTTTGATGCCAGCGGCTCTCGGATGGCATGGACGCTTATCGAGCAGCTTCAGGGTGGC	1452
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Qy	1658	atcaggttccaaggatccgaaccaccaaaagacaagaccatcatcctggagcagctgcgg	1717
Db	1513	GATAAATGGATTGGAGGGTCCCCCCCAGCTGACCAGACCCTGGTCATCAAGACATTCCGC	1572
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Db	1573		1632
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Db	1633	GCTGTTGTCTGTCCTTTAACATCTACAACTCACATGTCCGTTATATCCAGAACTCA	1692
Qу	1838	agtccatacatgaacaaccttatcatccttggagggatgctctcctatgcttccatattt	1897
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Qу	1898	ctctttggccttgatggatcctttgtctctgaaaagacctttgaaacactttgcaccgtc	1957
Db	1753	CCCCTGGGGCTCGATGGTTACCACATTGGGAGGAACCAGTTTCCTTTCGTCTGCCAGGCC	1812
QУ	1958	aggacctggattctcaccgtgggctacacgaccgcttttggggccatgtttgcaaagacc	2017
Db	1813	CGCCTCTGGCCTCGGGCCTGGGCTTTAGTCTGGGCTACGGTTCCATGTTCACCAAGATT	1872
Qу	2018	tggagagtccacgccatcttcaaaaatgtgaaaatgaagaagaagatcatcaaggacc	2076
Db	1873	TGGTGGGTCCACACGGTCTTCACAAAGAAGAAGAAGAAGAAGAAGAGAGTGGAGGAAGACTCTG	1932
Qу	2076	agaaactgcttgtgatcgtggggggcatgctgctgatcgacctgtgtatcctg	2128
Db	1933	GAACCCTGGAAGCTGTATGCCACAGTGGGCCTGCTGGTGGGCATGGATGTCCTCACTCTC	1992
Qу	2129	atctgctggcaggctgtggaccccctgcgaaggacagtggagaagtacagcatggagccg	2188
Db	1993	GCCATCTGGCAGATCGTGGACCCTCTGCACCGGACCATTGAGACATTTGCCAAGGAGGAA	2052
Qу	2189	gacccagcaggacgggatatctccatccgccctctcctggagcactgtgagaacacccat	2248
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Qу	2249	<pre>atgaccatctggcttggcatcgtctatgcctacaagggacttctcatgttgttcggttgt     </pre>	2308
Db	2113	ATGAATACATGGCTTGGCATTTTCTATGGTTACAAGGGGCTGCTGCTGCTGCTGGGAATC	2172
Qу	2309	ttcttagcttgggagacccgcaacgtcagcatccccgcactcaacgacagcaagtacatc	2368
Db	2173	TTCCTTGCTTATGAGACCAAGAGTGTGTCCACTGAGAAGATCAATGATCACCGGGCTGTG	2232
Qу	2369	gggatgagtgtctacaacgtggggatcatgtgcatcatcggggccgctgtctccttcct	2428
Db	2233	GGCATGGCTATCTACAATGTGGCAGTCCTGTGCCTCATCACTGCTCCTGTCACCATGATT	2292
Qу	2429	acccgggaccagcccaatgtgcagttctgcatcgtggctctggtcatcatcttctgcagc	2488
Db	2293	CTGTCCAGCCAGCAGGATGCAGCCTTTGCCTTTGCCTCTCTCCATAGTTTTCTCCTCC	2352
Qу	2489	accatcaccctctgcctggtattcgtgccgaagctcatcaccctgagaac 2538	
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V10266 standard; cDNA to mRNA; 2837 BP.
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AC
    03-JUN-1998 (first entry)
DT
    Rat GABA-BR1b receptor cDNA.
DF.
    Gamma-animobutyric acid; GABA-BR1b receptor; rat; brain; agonist;
KW
    inhibitory neurotransmitter; peripheral nervous system; antagonist;
KW
    treatment; dementia; depression; anxiety; bronchial inflammation; asthma;
KW
    epilepsy; cognitive function; ds.
KW
OS
    Rattus norvegicus.
                  Location/Qualifiers
FΗ
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FT
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FT
    WO9746675-A1.
PN
PD
    11-DEC-1997.
    19-MAR-1997; E01370.
ΡF
    22-NOV-1996; US-756091.
PR
    30-MAY-1996; US-655716.
PR
    (NOVS ) NOVARTIS AG.
PΑ
    Bettler B, Bittiger H, Froestl W, Kaupmann K, Mickel SJ;
PΙ
    WPI; 98-042183/04.
DR
    P-PSDB; W40118.
DR
    Purified GABA-B receptor or receptor protein - and antagonists of
PΤ
    these which may be useful in treating nervous system disorders
PT
    Claim 3; Page 67-74; 108pp; English.
PS
    This cDNA sequence encodes a novel rat GABA-B receptor protein,
CC
    GABA-BR1b. GABA (gamma-aminobutyric acid) is the major inhibitory
CC
    neurotransmitter found in the brain and peripheral nervous system
CC
    and this receptor may be used for the identification of {\sf GABA-B}
CC
    receptor agonists and antagonists. Such proteins may be used in
CC
    treatment of dementia, depression, anxiety, epilepsy, spasticity,
CC
    bronchial inflammation or asthma or to improve cognitive function.
CC
    GABA-B receptor ligands and probes derived from this sequence can be
CC
    used to assay for GABA-B receptors or DNA encoding them.
CC
                                          764 G;
                                 842 C;
             2837 BP;
                        621 A;
SO
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                       9.0%; Score 312.4; DB 1; Length 2837;
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Db
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Qу
                     301 CTCCGGTGTGGGCCTCTCACTCCCCTCATCTCCCGCGGCCTCACCCGAGGGTCCCCCCGC 360
Db
     438 cgccgcccagcagcccgccgctctccatcatgggcctcatgccgctcaccaaggaggtgg 497
Qу
                                              - 11
          361 ACCCCTCCTCAGAACGGCGTGCAGTATACATCGGGGCGCTGTTTCCCATGAGCGGGGGCT 420
Db
     498 ccaagggcagcatcgggcggtgtgctccccgccgtggaactggccatcgagcagatcc 557
Qу
                                                       1 1 11
                                       11
            | | | | |
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     421 GGCCGGGGGCCCAGCCTGCCAGCCCGCGTGGAGATGGCGCTGGAGGACGTTAACAGCC 480
Db
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Qу		gcaacgagtcactcctgcgccctacttcctcgacctgcggctctatgacacggagtgcg	
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Db		ACCCAGGGCAAGCCACCAAGTACTTGTACGAACTACTCTACAATGACCCCATCAAGATCA	
Qу		tggtgtttggaggcgtctgtccatccgtcacatccatcattgcagagtccctccaaggct	
Db		TTCTCATGCCTGGCTGTAGTTCTGTCTCCACACTTGTAGCTGAGGCTGCCCGGATGT	
Qу	738	ggaatctggtgcagctttcttttgctgcaaccacgcctgttctagccgataagaaaaaat	797
Db	655	GGAACCTTATTGTGCTCTCATATGGCTCCAGTTCACCAGCCTTGTCAAACCGACAGCGGT	714
Qу	798	accettatttettteggacegteecateagacaatgeggtgaateeageeattetgaagt	857
Db	715	TTCCCACGTTCTTCCGGACGCATCCATCCGCCACACTCCACAATCCCACCCGGGTGAAAC	774
Qу	858	tgctcaagcactaccagtggaagcgcgtgggcacgctgacgcaagacgttcagaggttct	917
Db	775		834
Qy	918	ctgaggtgcggaatgacctgactggagttctgtatggcgaggacattgagatttcagaca	977
Db	835	CCTCAACGCTGGATGACCTGGAGGAGCGAGTGAAAGAGGCTGGGATCGAGATCACTTTCC	894
Qу	978	ccgagagcttctccaacgatccctgtaccagtgtcaaaaagctgaaggggaatgatgtgc	1037
Db	895	GACAGAGTTTCTTCTCGGATCCAGCTGTGCCTGTTAAAAACCTGAAGCGTCAAGATGCTC	954
Qу	1038	ggatcatccttggccagtttgaccagaatatggcagcaaaagtgttctgttgtgcatacg	1097
Db	955	GAATCATCGTGGGACTTTTCTATGAGACGGAAGCCCGGAAAGTTTTTTGTGAGGTCTATA	1014
Qу	1098	aggagaacatgtatggtagtaaatatcagtggatcattccgggctggtacgagccttctt	1157
Db	1015	AGGAAAGGCTCTTTGGGAAGAAGTACGTCTGGTTCCTCATCGGGTGGTATGCTGACAACT	1074
Qу	1158	ggtgggagcaggtgcacacggaagccaactcatcccgctgcctccggaagaatctgcttg	1217
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Db	1126		1185
Qу	1278	ccatctcaggaaagactccacagcagtatgagagagtacaacaacaagcggtcagg	1336
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Qу	1336	cgtggggcccagcaagttccacgggtacgcctacgatggcatctgggtcatcgcca	1391
Db	1246	ACCCGAGGAGACTGGAGGCTTCCAGGAGGCACCACTGGCCTATGATGCTATCTGGGCCT	1305
Qу	1392	agacactgcagagggccatggagacactgcatgccagcagccggcaccagcggatccagg	1451
Db	1306		1365
Ov	1452	acttcaactacacqqaccacacqctqqqcaqqatcatcctcaatgccatgaacgagacca	1511

.

Db	1366		1425
Qу	1512	acttcttcggggtcacgggtcaagttgtattccggaatggggagagaatggggaccatta	1571
Db	1426	CCTTTGAGGGCGTTTCTGGCCATGTGGTCTTTGATGCCAGCGGCTCCCGGATGGCATGGA	1485
Qу	1572	aatttactcaatttcaagacagcagggaggtgaaggtgggagagtacaacgctgtggccg	1631
Db	1486	CACTTATCGAGCAGCTACAGGGCGGCAGCTACAAGAAGATCGGCTACTACGACAGCACCA	1545
Qу	1632	acacactggagatcatcaatgacaccatcaggttccaaggatccgaaccaccaaaagaca	1691
Db	1546	AGGATGATCTTTCCTGGTCCAAAACGGACAAGTGGATTGGAGGGTCTCCCCCAGCTGACC	1605
Qу		agaccatcatcctggagcagctgcggaagatctccctacctctctacagcatcctctctg	
Db	1606	AGACCTTGGTCATCAAGACATTCCGTTTCCTGTCTCAGAAACTCTTTATCTCCGTCTCAG	1665
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Db	1666	TTCTCTCCAGCCTGGGCATTGTTCTTGCTGTTGTCTGTCT	1725
Qу	1812	ggaatcagaagctcataaagatgtcgagtccatacatgaacaaccttatcatccttggag	1871
Db	1726	CCCACGTTCGTTATATCCAGAACTCCCAGCCCAACCTGAACAATCTGACTGTGGGCT	1785
Qу	1872	<pre>ggatgctctcctatgcttccatatttctctttggccttgatggatcctttgtctctgaaa  </pre>	1931
Db	1786	GCTCACTGGCACTGGCTGTCTTCCCTCTCGGGCTGGATGGTTACCACATAGGGAGAA	1845
Qу		agacctttgaaacactttgcaccgtcaggacctggattctcaccgtgggctacacgaccg	
Db	1846	GCCAGTTCCCGTTTGTCTGCCAGGCCCGCCTTTGGCTCTTGGGCTTTGGGCTTTAGTCTGG	1905
Qy	1992	cttttggggccatgtttgcaaagacctggagagtccacgccatcttcaaaaatgtgaaaa	2051
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Qу	2052	tgaagaagaagatcatcaaggaccagaaactgcttgtgatcgtggggggca	2102
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Qу		agggacttctcatgttgttcggttgtttcttagcttgggagacccgcaacgtcagcatcc	
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Qу	2463	tggctctggtcatcatcttctgcagcaccatcaccctctgcctggtattcgtgccgaagc	2522
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Qу	2523	tcatcaccctgagaac 2538	
Db	2446	TGCGCAGGCTGATCAC 2461	

```
V10264 standard; cDNA to mRNA; 4376 BP.
    V10264;
AC
    03-JUN-1998 (first entry)
DT
    Rat GABA-BR1a receptor cDNA.
DΕ
    Gamma-animobutyric acid; GABA-BR1a receptor; rat; brain; agonist;
KW
    inhibitory neurotransmitter; peripheral nervous system; antagonist;
KW
    treatment; dementia; depression; anxiety; bronchial inflammation; asthma;
ΚW
    epilepsy; cognitive function; ds.
ΚW
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OS
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FH
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                   /*tag= a
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                   /product= GABA-BR1a
FT
    WO9746675-A1.
PN
PD
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    19-MAR-1997; E01370.
ΡF
    22-NOV-1996; US-756091.
PR
    30-MAY-1996; US-655716.
PR
PΑ
    (NOVS ) NOVARTIS AG.
    Bettler B, Bittiger H, Froestl W, Kaupmann K, Mickel SJ;
PΙ
    WPI; 98-042183/04.
DR
    P-PSDB; W40116.
DR
    Purified GABA-B receptor or receptor protein - and antagonists of
PΤ
    these which may be useful in treating nervous system disorders
PΤ
    Claim 3; Page 42-50; 108pp; English.
PS
    This cDNA sequence encodes a novel rat GABA-B receptor protein,
CC
    GABA-BR1a. GABA (gamma-aminobutyric acid) is the major inhibitory
CC
    neurotransmitter found in the brain and peripheral nervous system
CC
    and this receptor may be used for the identification of GABA-B
CC
    receptor agonists and antagonists. Such proteins may be used in
CC
    treatment of dementia, depression, anxiety, epilepsy, spasticity,
CC
    bronchial inflammation or asthma or to improve cognitive function.
CC
    GABA-B receptor ligands and probes derived from this sequence can be
CC
    used to assay for GABA-B receptors or DNA encoding them.
CC
                                            1195 G;
                                                     1032 T;
                         940 A;
                                  1209 C;
             4376 BP;
SO
    Sequence
                        8.8%; Score 307.6; DB 1; Length 4376;
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                                                           27; Gaps
                                                                       4:
  Matches 968; Conservative
     570 tectgegecectaetteetegaeetgeggetetatgaeaeggagtgegaeaaegeaaaag 629
Qу
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Db
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Qу
          \perp
      969 TGCTCTCATATGGCTCCAGTTCACCAGCCTTGTCAAACCGACAGCGGTTTCCCACGTTCT 1028
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     810 ttcggaccgtcccatcagacaatgcggtgaatccagccattctgaagttgctcaagcact 869
Qу
                                              1 1111
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Qу		acggcagcaacaccagcggaccaccagggacggacggac	1169
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Qу		tgcacacggaagccaactcatcccgctgcctccggaagaatctgcttgct	
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Qу		cggaccacacgctgggcaggatcatcctcaatgccatgaacgagaccaacttcttcgggg	
Db	1680	ACAACCAGACCATTACAGACCAGATCTACCGGGCCATGAACTCCTCCTTTTGAGGGCG	1739
Qу	1524	tcacgggtcaagttgtattccggaatggggagagaatggggaccattaaatttactcaat	1583
Db	1740	TTTCTGGCCATGTGGTCTTTGATGCCAGCGGCTCCCGGATGGCATGGACACTTATCGAGC	1799
Qу	1584	ttcaagacagcagggaggtgaaggtgggagagtacaacgctgtggccgacacactggaga	1643
Db	1800	AGCTACAGGGCGGCAGCTACAAGAAGATCGGCTACTACGACAGCACCAAGGATGATCTTT	1859
Qy	1644	tcatcaatgacaccatcaggttccaaggatccgaaccaccaaaagacaagaccatcatcc	1703
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Db	2220	TGTTCACCAAGATCTGGTGGGTCCACACAGTCTTCACGAAGAAGGAGGAGAAGAAGGAGT	2279
Qу	2059	gaagatcatcaaggaccagaaactgcttgtgatcgtggggggcatgctgatcg	2114
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Db	2520	TGCTGCTGGGAATCTTTCTTGCTTACGAAACCAAGAGCGTGTCCACTGAAAAGATCAATG	2579
Qу	2355	acagcaagtacatcgggatgagtgtctacaacgtggggatcatgtgcatcatcggggccg	2414
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(FILE 'HOME' ENTERED AT 12:22:44 ON 01 JUN 1999)

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L1	6 S HG20
L2	6 DUP REM L1 (0 DUPLICATES REMOVED)
	E MCDONALD, T
	E MCDONALD T/AU
L3	152 S E3
	E MCDONALD TERRE/AU
L4	9 S E4-E6
	E BONNERT T/AU
L5	37 S E3-E7
L6	196 S L3 OR L4 OR L5
L7	22 S L6 AND (GABA? OR GAMMA?)
L8	11 DUP REM L7 (11 DUPLICATES REMOVED)
	FILE 'GENBANK' ENTERED AT 12:34:38 ON 01 JUN 1999
L9	2 S HG20

(FILE 'USPAT' ENTERED AT 12:13:09 ON 01 JUN 1999)
1 s HG20
72 s (GABA?(5A)RECEPTOR)/TI OR ((GABA?(5A)RECEPTOR?)/AB)

L1 L2

DUPLICATE 1 ANSWER 1 OF 11 MEDLINE 1.8

Stoichiometry of a ligand-gated ion channel determined by fluorescence energy transfer.

Farrar S J; Whiting P J; Bonnert T P; McKernan R M ΑU

JOURNAL OF BIOLOGICAL CHEMISTRY, (1999 Apr 9) 274 (15) 10100-4. Journal code: HIV. ISSN: 0021-9258.

We have developed a method to determine the stoichiometry of subunits within an oligomeric cell surface receptor using fluorescently tagged antibodies to the individual subunits and measuring energy transfer between them. Anti-c-Myc monoclonal antibody (mAb 9-E10) derivatized with a fluorophore (europium cryptate, EuK) was used to individually label c-Myc-tagged alphal-, beta2-, or gamma2-subunits of the hetero-oligomeric gamma-aminobutyric acid (GABAA) receptor in intact cells. The maximal fluorescent signal derived from the alpha1(c-Myc)beta2gamma2 and the alpha1beta2(c-Myc)gamma2 receptors was twice that obtained with alphalbeta2gamma2(c-Myc), suggesting that there are 2x alpha-, 2x beta-, and 1x gamma -subunits in a receptor monomer. This observation was extended using fluorescence energy transfer. Receptors were half-maximally saturated

with EuK-anti-c-Myc mAb, and the remaining alphal(c-Myc) subunits were labeled with excess anti-c-Myc mAb derivatized with the fluorescence energy acceptor, XL665. On exposure to laser light, energy transfer from EuK to XL665 occurred with alpha1(c-Myc)beta2gamma2 and alpha1beta2(c-Myc) gamma2, but no significant energy transfer was observed with alphalbeta2gamma2(c-Myc) receptors, indicating the absence of a second gamma-subunit in a receptor monomer. We confirm that the GABAA receptor subtype, alphalbeta2gamma2, is composed of two copies each of the alpha- and beta-subunits and one copy of the gamma-subunit (i.e. (alpha1)2(beta2)2(gamma2)1) and conclude that this method would have general applicability to other multisubunit cell surface proteins.

ANSWER 2 OF 11 MEDLINE L8

Identification of a GABAB receptor subunit, gb2, required for ΤI functional GABAB receptor activity.

Ng G Y; Clark J; Coulombe N; Ethier N; Hebert T E; Sullivan R; Kargman S; ΔIJ Chateauneuf A; Tsukamoto N; McDonald T; Whiting P; Mezey E; Johnson M P; Liu Q; Kolakowski L F Jr; Evans J F; Bonner T I; O'Neill G P SO

JOURNAL OF BIOLOGICAL CHEMISTRY, (1999 Mar 19) 274 (12) 7607-10.

Journal code: HIV. ISSN: 0021-9258. G protein-coupled receptors are commonly thought to bind their cognate AΒ ligands and elicit functional responses primarily as monomeric receptors. In studying the recombinant gamma-aminobutyric acid, type B ( GABAB) receptor (gbla) and a GABAB-like orphan receptor (gb2), we observed that both receptors are functionally inactive when expressed individually in multiple heterologous systems. Characterization of the tissue distribution of each of the receptors by in situ hybridization histochemistry in rat brain revealed co-localization of gbl and gb2 transcripts in many brain regions, suggesting the hypothesis that gb1 and gb2 may interact in vivo. In three established functional systems (inwardly rectifying K+ channel currents in Xenopus oocytes, melanophore pigment aggregation, and direct cAMP measurements in HEK-293 cells), GABA mediated a functional response in cells coexpressing gbla and qb2 but not in cells expressing either receptor individually. This GABA activity could be blocked with the GABAB receptor antagonist CGP71872. In COS-7 cells coexpressing gbla and gb2 receptors, co-immunoprecipitation of gbla and gb2 receptors was demonstrated,

indicating that abla and gb2 act as subunits in the formation of a functional GABA deceptor.

- L8 ANSWER 3 OF 11 BIOSIS COPYRIGHT 1999 BIOSIS DUPLICATE 2
- Cloning of a novel G-protein-coupled receptor GPR 51 resembling GABAB receptors expressed predominantly in nervous tissues and mapped proximal to the hereditary sensory neuropathy type 1 locus on chromosome 9.
- AU Ng, Gordon Y. K. (1); McDonald, Terrence; Bonnert, Tim; Rigby, Michael; Heavens, Robert; Whiting, Paul; Chateauneuf, Anne; Coulombe, Nathalie; Kargman, Stacia; Caskey, Thomas; Evans, Jilly; O'Neill, Gary P.; Liu, Qingyun
- SO Genomics, (March 15, 1999) Vol. 56, No. 3, pp. 288-295. ISSN: 0888-7543.

and

- AB Query of the expressed sequence tag database with the rat metabotropic GABABRIA receptor amino acid sequence using the TFASTA algorithm revealed two partial cDNA fragments whose sequence information was then used to isolate by PCR a novel full-length human cDNA encoding a putative G-protein-coupled receptor (GPCR), termed GPR 51. Sequence analysis revealed that it encoded a protein of 941 amino acids, similar in size
- homology to GABAB receptors followed by metabotropic glutamate receptors but not other GPCRs. GPR 51 expressed in COS-1 cells showed no specific binding for (3H)(+)baclofen and when expressed in Xenopus oocyte and Xenopus melanophore functional assays showed no activity to GABA, (-)baclofen, and glutamic acid. Northern blot analysis and in situ hybridization revealed that GPR 51 transcripts were predominantly expressed in the central nervous system with highest abundance in the cortex, thalamus, hippocampus, amygdala, cerebellum, and spinal cord. In contrast, GPR 51 receptor transcripts were almost not detected in the peripheral tissues. Gene GPR 51 was localized by radiation hybrid mapping to chromosome 9, 4.81 cR from the WI-8684 marker, and proximal to the hereditary sensory neuropathy type 1 locus.
- L8 ANSWER 8 OF 11 MEDLINE DUPLICATE 6
- TI The use of expressed sequence tag databases to identify novel human gamma-aminobutyric acid type receptor genes.
- AU Whiting P J; McAllister G; Bonnert T; Heavens R P; Rigby M R; Sirinathsinghji D J; Marshall G; Thompson S A; Wafford K A
- SO BIOCHEMICAL SOCIETY TRANSACTIONS, (1997 Aug) 25 (3) 817-9. Ref: 21 Journal code: E48. ISSN: 0300-5127.